

165 170 175
 Lys Lys Glu Asn Gly Leu Val Ile Leu Thr Ser His Ile Ser Glu Asp
 180 185 190
 Ile Ser Asp Leu Cys Thr Asp Val Leu Val Val Glu Asn Gly His Ile
 195 200 205
 Gln Met
 210
 <210> 365
 <211> 97
 <212> PRT
 <213> Streptococcus pneumoniae
 <400> 365
 Met Arg Ser Met Thr Arg Leu Ala Ser Gln Val Ser Ser Phe Pro Cys
 1 5 10 15
 Phe Thr Phe Ala Lys Cys Phe Ser Lys Ser Ser Lys Leu Lys Leu Asp
 20 25 30
 Val Lys Lys Val Gly Lys Phe Ser Cys Ile Arg Tyr Trp Arg Met Thr
 35 40 45
 Cys Arg Ile Ser Ser Arg Thr Gln Thr Val Asp Cys Ser Ala Pro Ile
 50 55 60
 Ser Ser Lys Ile Arg Thr Ser Asp Ser Leu Ile Ser Ser Thr Lys Val
 65 70 75 80
 Leu Thr Leu Pro Ser Leu Ile Ala Phe Leu Thr Ser Met Thr Lys Leu
 85 90 95
 Gly

<210> 366
 <211> 501
 <212> PRT
 <213> Streptococcus pneumoniae

<400> 366

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Met Ile Ile Leu Lys Glu Phe Lys Pro Met Ser Leu Leu Val Phe Glu
 1           5           10           15

Asn Val Ser Lys Ser Tyr Gly Ala Thr Pro Ala Leu Glu Asn Val Ser
          20           25           30

Leu Asp Ile Pro Ala Gly Lys Ile Val Gly Leu Leu Gly Pro Asn Gly
 35           40           45

Ser Gly Lys Thr Thr Leu Ile Lys Leu Ile Asn Gly Leu Leu Gln Pro
 50           55           60

Asp Gln Gly Arg Val Leu Ile Asn Asp Met Asp Pro Ser Pro Ala Thr
 65           70           75           80

Lys Ala Val Val Ala Tyr Leu Pro Asp Thr Thr Tyr Leu Asn Glu Gln
          85           90           95

Met Lys Val Lys Glu Ala Leu Thr Tyr Phe Lys Thr Phe Tyr Lys Asp
          100          105          110

Phe Asn Leu Glu Arg Ala His His Leu Leu Ala Asp Leu Gly Ile Asp
 115           120           125

Glu Asn Ser Arg Leu Lys Lys Leu Ser Lys Glu Asn Lys Glu Lys Val
 130           135           140

Gln Leu Ile Leu Val Met Ser Arg Asp Ala Arg Leu Tyr Val Leu Asp
 145           150           155           160

Glu Pro Ile Gly Gly Val Asp Pro Ala Ala Arg Ala Tyr Ile Leu Asn
          165           170           175

Thr Ile Ile Asn Asn Tyr Ser Pro Thr Ser Thr Val Leu Ile Ser Thr
          180          185           190

His Leu Ile Ser Asp Ile Glu Pro Ile Leu Asp Glu Ile Val Phe Leu
 195           200           205

Lys Asp Gly Lys Val Val Arg Gln Gly Asn Val Asp Asp Ile Arg Tyr

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210	215	220
Glu Ser Gly Glu Ser Ile Asp Gln Leu Phe Arg Gln Asn Leu Arg Pro		
225	230	235 240
Lys Gln Arg Arg Leu Phe Met Phe Trp Asn Leu Val Arg Tyr Glu Phe		
	245	250 255
Lys Asn Val Asn Lys Trp Tyr Leu Ala Leu Tyr Ala Ala Val Leu Val		
	260	265 270
Leu Ser Ala Leu Ile Gly Ile Gln Thr Gln Gly Phe Lys Asn Leu Pro		
	275	280 285
Tyr Gln Glu Ser Gln Ala Thr Met Leu Leu Phe Leu Ala Thr Val Phe		
	290	295 300
Gly Gly Leu Met Leu Thr Leu Gly Ile Ser Thr Ile Phe Leu Ile Ile		
	305	310 315 320
Lys Arg Phe Lys Gly Ser Val Tyr Asp Arg Gln Gly Tyr Leu Thr Leu		
	325	330 335
Thr Leu Pro Val Ser Glu His His Ile Ile Thr Ala Lys Leu Ile Gly		
	340	345 350
Ala Phe Ile Trp Ser Leu Ile Ser Thr Ala Val Leu Ala Leu Ser Ala		
	355	360 365
Val Ile Ile Leu Ala Leu Thr Ala Pro Glu Trp Ile Pro Leu Ser Tyr		
	370	375 380
Val Ile Thr Phe Val Glu Thr His Leu Pro Gln Ile Phe Leu Thr Gly		
	385	390 395 400
Ile Ser Phe Leu Leu Asn Thr Ile Ser Gly Ile Leu Cys Ile Tyr Leu		
	405	410 415
Ala Ile Ser Ile Gly Gln Leu Phe Asn Glu Tyr Arg Thr Ala Leu Ala		
	420	425 430

Val Ala Val Tyr Ile Gly Ile Gln Ile Val Ile Gly Phe Ile Glu Leu
 435 440 445

Phe Phe Asn Leu Ser Ser Asn Phe Tyr Val Asn Ser Leu Val Gly Leu
 450 455 460

Asn Asp His Phe Tyr Met Gly Ala Gly Ile Ala Ile Val Glu Glu Leu
 465 470 475 480

Ile Phe Ile Ala Ile Phe Tyr Leu Gly Thr Tyr Tyr Ile Leu Arg Asn
 485 490 495

Lys Val Asn Leu Leu
 500

<210> 367

<211> 381

<212> PRT

<213> Streptococcus pneumoniae

<400> 367

Met Val Thr Pro Phe Val Thr Leu Leu Val Met Ser Ile Leu Gly Leu
 1 5 10 15

Phe Val Ile Gly Pro Val Phe His Val Val Glu Asn Tyr Ile Leu Ile
 20 25 30

Ala Thr Lys Ala Ile Leu Ser Met Pro Phe Gly Leu Gly Gly Phe Leu
 35 40 45

Ile Gly Gly Val His Gln Leu Ile Val Val Ser Gly Val His His Ile
 50 55 60

Phe Asn Leu Leu Glu Val Gln Leu Leu Ala Ala Asp His Ala Asn Pro
 65 70 75 80

Phe Asn Ala Ile Ile Thr Ala Ala Met Thr Ala Gln Gly Ala Ala Thr
 85 90 95

Val Ala Val Gly Val Lys Thr Lys Asn Pro Lys Leu Lys Thr Leu Ala
 100 105 110

Phe Pro Ala Ala Leu Ser Ala Phe Leu Gly Ile Thr Glu Pro Ala Ile
 115 120 125

Phe Gly Val Asn Leu Arg Phe Arg Lys Pro Phe Phe Leu Ser Leu Ile
 130 135 140

Ala Gly Ala Ile Gly Gly Gly Leu Ala Ser Ile Leu Gly Leu Ala Gly
 145 150 155 160

Thr Gly Asn Gly Ile Thr Ile Ile Pro Gly Thr Met Leu Tyr Val Gly
 165 170 175

Asn Gly Gln Leu Pro Gln Tyr Leu Leu Met Val Ala Val Ser Phe Ala
 180 185 190

Leu Gly Phe Ala Leu Thr Tyr Met Phe Gly Tyr Glu Asp Glu Val Asp
 195 200 205

Ala Thr Ala Ala Ala Lys Arg Ala Glu Val Ala Glu Glu Lys Glu Glu
 210 215 220

Val Ala Pro Ala Ala Leu Gln Asn Glu Thr Leu Val Thr Pro Ile Val
 225 230 235 240

Gly Asp Val Val Ala Leu Ala Asp Val Asn Asp Pro Val Phe Ser Ser
 245 250 255

Gly Ala Met Gly Gln Gly Ile Val Val Lys Pro Ser Gln Gly Val Val
 260 265 270

Tyr Ala Pro Ala Asp Ala Glu Val Ser Ile Ala Phe Pro Thr Gly His
 275 280 285

Ala Phe Gly Leu Lys Thr Arg Asn Gly Ala Glu Val Leu Ile His Val
 290 295 300

Gly Ile Asp Thr Val Ser Met Asn Gly Asp Gly Phe Glu Thr Lys Val
 305 310 315 320

Ala Gln Gly Asn Lys Val Lys Ala Gly Asp Val Leu Gly Thr Phe Asp
 325 330 335

Ser Asn Lys Ile Ala Ala Ala Gly Leu Asp Asp Thr Thr Met Val Ile
 340 345 350

Val Thr Asn Thr Gly Asp Tyr Ala Ser Val Ala Pro Val Ala Thr Gly
 355 360 365

Ser Val Ala Lys Gly Asp Ala Val Ile Glu Val Lys Ile
 370 375 380

<210> 368

<211> 249

<212> PRT

<213> Streptococcus pneumoniae

<400> 368

Met Gln Met Asn Asn Gln Glu Ile Ala Lys Lys Val Ile Asp Ala Leu
 1 5 10 15

Gly Gly Arg Glu Asn Val Asn Ser Val Ala His Cys Ala Thr Arg Leu
 20 25 30

Arg Val Met Val Lys Asp Glu Glu Lys Ile Asn Lys Glu Val Ile Glu
 35 40 45

Asn Leu Glu Lys Val Gln Gly Ala Phe Phe Asn Ser Gly Gln Tyr Gln
 50 55 60

Ile Ile Phe Gly Thr Gly Thr Val Asn Lys Met Tyr Asp Glu Val Val
 65 70 75 80

Val Leu Gly Leu Pro Thr Ser Ser Lys Asp Asp Met Lys Ala Glu Val
 85 90 95

Ala Lys Gln Gly Asn Trp Phe Gln Arg Ala Ile Arg Thr Phe Gly Asp
 100 105 110

Val Phe Val Pro Ile Ile Pro Val Ile Val Ala Thr Gly Leu Phe Met
 115 120 125

Gly Val Arg Gly Leu Phe Asn Ala Leu Glu Met Pro Leu Pro Gly Asp
 130 135 140

Phe Ala Thr Tyr Thr Gln Ile Leu Thr Asp Thr Ala Phe Ile Ile Leu
145 150 155 160

Pro Gly Leu Val Val Trp Ser Thr Phe Arg Val Phe Gly Gly Asn Pro
165 170 175

Ala Val Gly Ile Val Leu Leu Gly Met Met Leu Val Ser Gly Ser Leu Pro
180 185 190

Asn Ala Trp Ala Val Ala Gln Gly Gly Glu Val Thr Ala Met Asn Phe
195 200 205

Phe Gly Phe Ile Pro Val Val Gly Leu Gln Gly Ser Val Leu Pro Ala
210 215 220

Phe Ile Ile Gly Val Val Gly Ala Lys Phe Glu Lys Ala Val Arg Lys
225 230 235 240

Leu Phe Gln Met Ser Leu Thr Ser Trp
245

<210> 369

<211> 201

<212> PRT

<213> Streptococcus pneumoniae

<400> 369

Met Arg Leu Met Lys Ile Trp Tyr Asn Glu Ile Lys Glu Phe Tyr Met
1 5 10 15

Gln Lys Phe Ile Gln Ala Tyr Ile Glu Lys Leu Asp Val Thr Thr Ile
20 25 30

Ile Glu Asn Ile Leu Thr Lys Val Ile Ser Leu Leu Leu Leu Ile
35 40 45

Val Phe Tyr Ile Ala Lys Lys Met Leu His Thr Met Val Gln Arg Ile
50 55 60

Val Lys Pro Ser Leu Lys Met Ser Arg His Asp Val Gly Arg Gln Lys
65 70 75 80

Thr Ile Ser Arg Leu Leu Glu Asn Val Phe Asn Tyr Thr Leu Tyr Phe
85 90 95

Phe Leu Leu Tyr Cys Ile Leu Ser Ile Leu Gly Leu Pro Val Ser Ser
100 105 110

Leu Leu Ala Gly Ala Gly Ile Ala Gly Val Ala Ile Gly Met Gly Ala
115 120 125

Gln Gly Phe Leu Ser Asp Val Ile Asn Gly Phe Phe Ile Leu Phe Glu
130 135 140

Arg Gln Leu Asp Val Gly Asp Glu Val Val Leu Thr Asn Gly Pro Ile
145 150 155 160

Thr Val Ser Gly Lys Val Val Ser Val Gly Ile Arg Thr Thr Gln Leu
165 170 175

Arg Ser Glu Glu Gln Ala Leu His Phe Val Pro Asn Arg Asn Ile Thr
180 185 190

Val Val Ser Asn Phe Ser Arg Thr Asp
195 200

<210> 370

<211> 598

<212> PRT

<213> Streptococcus pneumoniae

<400> 370

Met Glu Gln Lys His Arg Ser Glu Phe Pro Glu Lys Glu Leu Trp Asp
1 5 10 15

Leu Thr Ala Leu Tyr Gln Asp Arg Glu Asp Phe Leu Arg Ala Ile Glu
20 25 30

Lys Ala Arg Glu Asp Ile Asn Gln Phe Ser Arg Asp Tyr Lys Gly Asn
35 40 45

Leu His Thr Phe Glu Asp Phe Glu Lys Ala Phe Ala Glu Leu Glu Gln
50 55 60

Ile Tyr Ile Gln Met Ser His Ile Gly Asn Tyr Gly Phe Met Pro Gln
 65 70 75 80

Thr Thr Asp Tyr Ser Asn Asp Glu Phe Ala Asn Ile Ala Gln Ala Gly
 85 90 95

Met Glu Phe Glu Thr Asp Ala Ser Val Ala Leu Thr Phe Phe Asp Asp
 100 105 110

Ala Leu Val Ala Ala Asp Glu Glu Val Leu Asp Arg Leu Gly Lys Leu
 115 120 125

Pro His Leu Thr Ala Ala Ile Arg Gln Ala Lys Ile Lys Lys Ala His
 130 135 140

Tyr Leu Gly Ala Asp Val Glu Lys Ala Leu Thr Asn Leu Gly Glu Val
 145 150 155 160

Phe Tyr Ser Pro Gln Asp Ile Tyr Thr Lys Met Arg Ala Gly Asp Phe
 165 170 175

Glu Met Ala Asp Phe Glu Ala His Gly Lys Thr Tyr Lys Asn Ser Phe
 180 185 190

Val Thr Tyr Glu Asn Phe Tyr Gln Asn His Glu Asp Ala Glu Val Arg
 195 200 205

Glu Lys Ser Phe Arg Ser Phe Ser Glu Gly Leu Arg Lys His Gln Asn
 210 215 220

Thr Ala Ala Ala Ala Tyr Leu Ala Gln Val Lys Ser Glu Lys Leu Leu
 225 230 235 240

Ala Asp Met Lys Gly Tyr Asp Ser Val Phe Asp Tyr Leu Leu Ala Glu
 245 250 255

Gln Glu Val Asp Arg Val Met Phe Asp Arg Gln Ile Asp Leu Ile Met
 260 265 270

Lys Asp Phe Ala Pro Val Ala Gln Arg Tyr Leu Lys His Val Ala Lys
 275 280 285

Val Asn Gly Leu Glu Lys Met Thr Phe Ala Asp Trp Lys Leu Asp Leu
 290 295 300

Asp Ser Ala Leu Asn Pro Glu Val Thr Ile Asp Asp Ala Tyr Asp Leu
 305 310 315 320

Val Met Lys Ser Val Glu Pro Leu Gly Gln Glu Tyr Cys Gln Glu Val
 325 330 335

Ala Arg Tyr Gln Glu Glu Arg Trp Val Asp Phe Ala Ala Asn Ser Gly
 340 345 350

Lys Asp Ser Gly Gly Tyr Ala Ala Asp Pro Tyr Arg Val His Pro Tyr
 355 360 365

Val Leu Met Ser Trp Thr Gly Arg Leu Ser Asp Val Tyr Thr Leu Ile
 370 375 380

His Glu Ile Gly His Ser Gly Gln Phe Ile Phe Ser Asp Asn His Gln
 385 390 395 400

Ser Tyr Phe Asn Ala His Met Ser Thr Tyr Tyr Val Glu Ala Pro Ser
 405 410 415

Thr Phe Asn Glu Leu Leu Leu Ser Asp Tyr Leu Glu Asn Gln Ser Asn
 420 425 430

Asp Pro Arg Gln Lys Arg Phe Ala Leu Ala His Arg Leu Thr Asp Thr
 435 440 445

Tyr Phe His Asn Phe Ile Thr His Leu Leu Glu Ala Ala Phe Gln Arg
 450 455 460

Lys Val Tyr Thr Leu Ile Glu Glu Gly Glu Thr Phe Gly Ala Ser Lys
 465 470 475 480

Leu Asn Ser Ile Met Lys Glu Val Leu Thr Asp Phe Trp Gly Asp Ala
 485 490 495

Ile Glu Ile Asp Asp Asp Ala Thr Leu Thr Trp Met Arg Gln Ala His

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500                                505                                510

Tyr Tyr Met Gly Leu Tyr Ser Tyr Thr Tyr Ser Ala Gly Leu Val Ile
515                                520                                525

Ser Thr Ala Gly Tyr Leu His Leu Lys His Ser Glu Thr Gly Ala Glu
530                                535                                540

Asp Trp Leu Asn Leu Leu Lys Ser Gly Gly Ser Lys Thr Pro Leu Glu
545                                550                                555                                560

Ser Ala Met Ile Ile Gly Ala Asp Ile Ser Thr Asp Lys Pro Leu Arg
565                                570                                575

Asp Thr Ile Gln Phe Leu Ser Asp Thr Val Asp Gln Ile Ile Ser Tyr
580                                585                                590

Ser Ala Glu Leu Gly Glu
595

<210> 371
<211> 190
<212> PRT
<213> Streptococcus pneumoniae

<400> 371

Met Tyr Met Ser Lys Ala Lys Lys Ile Cys Phe Ile Ile Phe Cys Ile
1 5 10 15

Leu Ile Leu Thr Ile Phe Leu Pro Val Leu Ile Asp Tyr His Gln Val
20 25 30

Ser Asp Leu Gly Ile His Leu Leu Ser Trp Arg Gln Asn Ser Val Val
35 40 45

Glu Phe Tyr Leu Ala Arg Tyr Val Phe Trp Gly Thr Val Val Leu Ser
50 55 60

Thr Leu Val Leu Leu Ser Ile Leu Val Val Met Phe Tyr Pro Lys Arg
65 70 75 80

Tyr Leu Glu Ile Gln Leu Glu Thr Lys Asn Asp Thr Leu Lys Leu Lys

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	85	90	95
Asn Ser Ala Ile Glu Gly Phe Val Arg Ser Leu Val Ser Asp His Arg	100	105	110
Leu Ile Lys Asn Pro Thr Val His Val Asn Leu Arg Lys Asn Lys Cys	115	120	125
Phe Val His Val Glu Gly Lys Ile Leu Pro Ser Asp Asn Ile Ala Asp	130	135	140
Arg Cys Gln Ile Ile Gln Asn Glu Ile Thr Asn Gly Leu Lys Gln Phe	145	150	155
Phe Gly Ile Glu Arg Gln Val Lys Leu Glu Val Ala Val Lys Asn Tyr	165	170	175
Gln Pro Lys Pro Gln Asn Lys Lys Thr Val Ser Arg Val Lys	180	185	190
<210> 372			
<211> 219			
<212> PRT			
<213> Streptococcus pneumoniae			
<400> 372			
Met Ile Asp Phe Tyr Phe Phe Leu Val Gly Ser Ile Leu Ala Ser Phe	1	5	10
Leu Gly Leu Val Ile Asp Arg Phe Pro Glu Gln Ser Ile Ile Ser Ser	20	25	30
Ala Ser His Cys Asp Ser Cys Gln Thr Pro Leu Arg Pro Leu Asp Leu	35	40	45
Ile Pro Ile Leu Ser Gln Val Phe Asn Arg Phe Arg Cys Arg Tyr Cys	50	55	60
Lys Val Arg Tyr Pro Val Trp Tyr Ala Leu Phe Glu Leu Ser Leu Gly	65	70	75
Leu Leu Phe Leu Leu Tyr Ser Trp Gly Trp Leu Ser Leu Gly Gln Val			80

	85	90	95
Val Leu Ile Thr Ala Gly Leu Thr Leu Gly Ile Tyr Asp Phe His His	100	105	110
Gln Glu Tyr Pro Leu Leu Val Trp Met Thr Phe Gln Leu Ile Leu Ile	115	120	125
Ala Ser Ser Gly Trp Asn Leu Val Met Val Ser Phe Leu Ile Leu Gly	130	135	140
Ile Leu Ala His Phe Ile Asp Ile Arg Met Gly Ala Gly Asp Phe Leu	145	150	155
Phe Leu Ala Ser Cys Ala Leu Val Phe Ser Val Thr Glu Leu Leu Ile	165	170	175
Leu Ile Gln Phe Ala Ser Ala Thr Gly Ile Leu Ala Phe Leu Leu Gln	180	185	190
Lys Lys Lys Glu Arg Leu Pro Phe Val Pro Phe Leu Leu Leu Ala Thr	195	200	205
Cys Leu Ile Ile Phe Gly Lys Leu Leu Leu Val	210	215	
<210> 373			
<211> 383			
<212> PRT			
<213> Streptococcus pneumoniae			
<400> 373			
Met Phe Asn Gly Arg Val Leu Lys Glu Leu Arg Leu Leu Asn Gly Leu	1	5	10
Ser Arg Ala Glu Leu Ala Gln Arg Ile Asn Leu Thr Glu Gln Ala Ile	20	25	30
Trp Gln Phe Glu Ser Asn Glu Thr Lys Pro Lys Leu Ser Thr Lys Met	35	40	45
His Leu Ala Asn Gln Phe His Val Asp Leu Thr Tyr Phe Glu Gln Glu			

50	55	60
Glu Glu Ser Ile Arg Phe Asp Ser Ser Val Ile Ala Phe Arg Asn Ala .		
65	70	75 80
Asp Leu Ala Thr Arg Lys Thr Ile Asp Ile Gln Thr Met Tyr Leu His		
	85	90 95
Lys Val Asp Ser Leu Ile Asp Tyr Phe Glu Ser Phe Val Ile Ile Pro		
	100	105 110
Asn Ile Ile Ile His Asp Leu Ser Asn Val Val Ser Glu Ser Tyr His		
	115	120 125
Lys Gly Glu Ser Ile Glu Glu Leu Ala Leu Tyr Ala Arg Glu Lys Leu		
	130	135 140
Gly Ile Ser Lys Asp Asn His Asp Leu Leu Tyr Lys Leu Glu Arg Ser		
	145	150 155 160
Gly Ile Tyr Ile Val Glu Arg Leu Ile Asn Gly Gln Ala Asp Ala Tyr		
	165	170 175
Ser Ala Trp Ser Lys Leu Gly Arg Pro Tyr Ile Val Leu Gly Thr Asn		
	180	185 190
Lys Ser Ser Val Arg Arg Asn Phe Asp Leu Ala His Glu Leu Gly His		
	195	200 205
Ile Leu Leu His Lys Tyr Lys Asp Met Asn Glu Asp Gly Asp Arg Leu		
	210	215 220
Glu Gln Glu Ala Asn Tyr Phe Ala Ser Cys Phe Leu Leu Pro Lys Glu		
	225	230 235 240
Glu Phe Leu Val Lys Phe Glu Glu Arg Val Gly Lys Arg Val Ser Asn		
	245	250 255
Pro Asp Ser Tyr Ile Leu Leu Lys Ser Asp Leu Asn Val Ser Ile Gln		
	260	265 270

Ala Leu Glu Tyr Arg Ala Phe Lys Leu Gly Leu Leu Thr Pro Lys Gln
 275 280 285

His Ser Tyr Phe Tyr Arg Gln Ile Ala Gln Lys Gly Tyr Lys Met Ile
 290 295 300

Glu Pro Leu Asp Asp Gln Ile Phe Val Lys Lys Pro Ser Lys Val Lys
 305 310 315 320

Ser Ile Leu Asp Val Val Leu Ser Asn His Leu Val Ser Leu Ala Thr
 325 330 335

Ile Met Ser Lys Gln Ser Ile Arg Leu Gln Phe Ile Ser Glu Ile Phe
 340 345 350

Ser Val Glu Met Lys Phe Phe Asp Gln Tyr Gln Glu Asp Arg Arg Thr
 355 360 365

Asp Arg Phe Asp Asn Ile Ile Pro Leu Tyr Lys Arg Asn Asn Leu
 370 375 380

<210> 374

<211> 594

<212> PRT

<213> Streptococcus pneumoniae

<400> 374

Met Ala Ser Gly Phe His Leu Gln Ser Glu Arg Asp Phe Met Ser Ile
 1 5 10 15

Ile Gln Lys Leu Trp Trp Phe Phe Lys Leu Glu Lys Arg Arg Tyr Leu
 20 25 30

Val Gly Ile Val Ala Leu Ile Leu Val Ser Val Leu Asn Leu Ile Pro
 35 40 45

Pro Met Val Met Gly Arg Val Ile Asp Ala Ile Thr Ser Gly Gln Leu
 50 55 60

Thr Gln Gln Asp Leu Leu Leu Ser Leu Phe Tyr Leu Leu Leu Ala Ala
 65 70 75 80

Phe Gly Met Tyr Tyr Leu Arg Tyr Val Trp Arg Met Tyr Ile Leu Gly
 85 90 95

Thr Ser Tyr Cys Leu Gly Gln Ile Met Arg Ser Arg Leu Phe Lys His
 100 105 110

Phe Thr Lys Met Ser Ser Ala Phe Tyr Gln Thr Tyr Arg Thr Gly Asp
 115 120 125

Leu Met Ala His Ala Thr Asn Asp Ile Asn Ala Leu Thr Arg Leu Ala
 130 135 140

Gly Gly Gly Val Met Ser Ala Val Asp Ala Ser Ile Thr Ala Leu Val
 145 150 155 160

Thr Leu Leu Thr Met Leu Phe Ser Ile Ser Trp Gln Met Thr Leu Val
 165 170 175

Ala Ile Leu Pro Leu Pro Phe Met Ala Tyr Thr Thr Ser Arg Leu Gly
 180 185 190

Arg Lys Thr His Lys Ala Phe Gly Glu Ser Gln Ala Ala Phe Ser Glu
 195 200 205

Leu Asn Asn Lys Val Gln Glu Ser Val Ser Gly Ile Lys Val Thr Lys
 210 215 220

Ser Phe Gly Tyr Gln Ala Asp Glu Leu Lys Ser Phe Gln Ala Val Asn
 225 230 235 240

Glu Leu Thr Phe Gln Lys Asn Leu Gln Thr Met Lys Tyr Asp Ser Leu
 245 250 255

Phe Asp Pro Met Val Leu Leu Phe Val Gly Ser Ser Tyr Val Leu Thr
 260 265 270

Leu Leu Val Gly Ser Leu Met Val Gln Glu Gly Gln Ile Thr Val Gly
 275 280 285

Asn Leu Val Thr Phe Ile Ser Tyr Leu Asp Met Leu Val Trp Pro Leu
 290 295 300

Leu Ala Ile Gly Phe Leu Phe Asn Thr Thr Gln Arg Gly Lys Val Ser
 305 310 315 320

Tyr Gln Arg Ile Glu Asn Leu Leu Ser Gln Glu Ser Pro Val Gln Asp
 325 330 335

Pro Glu Phe Pro Leu Asp Gly Ile Glu Asn Gly Arg Leu Glu Tyr Ala
 340 345 350

Ile Asp Ser Phe Ala Phe Glu Asn Glu Glu Thr Leu Thr Asp Ile His
 355 360 365

Phe Ser Leu Ala Lys Gly Gln Thr Leu Gly Leu Val Gly Gln Thr Gly
 370 375 380

Ser Gly Lys Thr Ser Leu Ile Lys Leu Leu Leu Arg Glu Tyr Asp Val
 385 390 395 400

Asp Lys Gly Ala Ile Tyr Leu Asn Gly His Asp Ile Arg Asp Tyr Arg
 405 410 415

Leu Thr Asp Leu Arg Ser Leu Met Gly Tyr Val Pro Gln Asp Gln Phe
 420 425 430

Leu Phe Ala Thr Ser Ile Leu Asp Asn Ile Arg Phe Gly Asn Pro Asn
 435 440 445

Leu Pro Leu Ser Ala Val Glu Glu Ala Thr Lys Leu Ala Arg Val Tyr
 450 455 460

Gln Asp Ile Val Asp Met Pro Gln Gly Phe Asp Thr Leu Ile Gly Glu
 465 470 475 480

Lys Gly Val Ser Leu Ser Gly Gly Gln Lys Gln Arg Leu Ala Met Ser
 485 490 495

Arg Ala Met Ile Leu Asp Pro Asp Ile Leu Ile Leu Asp Asp Ser Leu
 500 505 510

Ser Ala Val Asp Ala Lys Thr Glu Tyr Ala Ile Ile Asp Asn Leu Lys
 515 520 525

Glu Met Arg Lys Asp Lys Thr Thr Ile Ile Thr Ala His Arg Leu Ser
 530 535 540

Ala Val Val His Ala Asp Phe Ile Leu Val Leu Gln Asn Gly Gln Ile
 545 550 555 560

Ile Glu Arg Gly Thr His Glu Asp Leu Leu Ala Leu Asp Gly Trp Tyr
 565 570 575

Ala Gln Thr Tyr Gln Ser Gln Gln Leu Glu Met Lys Gly Glu Glu Asp
 580 585 590

Ala Glu

<210> 375

<211> 655

<212> PRT

<213> Streptococcus pneumoniae

<400> 375

Met Gly Lys Phe Glu Gln Glu Ala Lys Asp Leu Leu Gln Ala Ile Gly
 1 5 10 15

Gly Lys Glu Asn Val Thr Ala Val Thr His Cys Ala Thr Arg Met Arg
 20 25 30

Phe Val Leu Gly Asp Asp Lys Lys Ala Asn Val Lys Ala Ile Glu Ser
 35 40 45

Ile Pro Ala Val Lys Gly Thr Phe Thr Asn Ala Gly Gln Phe Gln Val
 50 55 60

Ile Ile Gly Asn Asp Val Pro Ile Phe Tyr Asn Asp Phe Thr Ala Val
 65 70 75 80

Ser Gly Ile Glu Gly Val Ser Lys Glu Ala Ala Lys Ser Ala Ala Lys
 85 90 95

Ser Asn Gln Asn Val Val Gln Gly Val Met Thr Thr Leu Ala Glu Ile
 100 105 110

Phe Thr Pro Ile Ile Pro Ala Leu Ile Val Gly Gly Leu Ile Leu Gly
 115 120 125

Phe Arg Asn Val Leu Glu Gly Val His Trp Ser Met Leu Asp Gly Lys
 130 135 140

Thr Ile Thr Glu Ser Ser Gln Phe Trp Ala Gly Val Asn His Phe Leu
 145 150 155 160

Trp Leu Pro Gly Glu Ala Ile Phe Gln Phe Leu Pro Val Gly Ile Thr
 165 170 175

Trp Ser Val Ser Arg Lys Met Gly Thr Ser Gln Ile Leu Gly Ile Val
 180 185 190

Leu Gly Ile Cys Leu Val Ser Pro Gln Leu Leu Asn Ala Tyr Ala Val
 195 200 205

Ala Ser Thr Pro Ala Ala Asp Ile Ala Ala Asn Trp Val Trp Asn Phe
 210 215 220

Gly Tyr Phe Thr Val Asn Arg Ile Gly Tyr Gln Ala Gln Val Ile Pro
 225 230 235 240

Ala Leu Leu Ala Gly Leu Ser Leu Ser Tyr Leu Glu Ile Phe Trp His
 245 250 255

Lys His Ile Pro Glu Val Ile Ser Met Ile Phe Val Pro Phe Leu Ser
 260 265 270

Leu Ile Pro Ala Leu Ile Leu Ala His Thr Val Leu Gly Pro Ile Gly
 275 280 285

Trp Thr Ile Gly Gln Gly Leu Ser Ser Val Val Leu Ala Gly Leu Thr
 290 295 300

Gly Pro Val Lys Trp Leu Phe Gly Ala Ile Phe Gly Ala Leu Tyr Ala
 305 310 315 320

Pro Phe Val Ile Thr Gly Leu His His Met Thr Asn Ala Ile Asp Thr

325	330	335
Gln Leu Ile Ala Asp Ala Gly Gly Thr Ala Leu Trp Pro Met Ile Ala 340 345 350		
Leu Ser Asn Ile Ala Gln Gly Ser Ala Val Phe Ala Tyr Tyr Phe Met 355 360 365		
His Arg His Asp Glu Arg Glu Ala Gln Val Ser Leu Pro Ala Thr Ile 370 375 380		
Ser Ala Tyr Leu Gly Val Thr Glu Pro Ala Leu Phe Gly Val Asn Val 385 390 395 400		
Lys Tyr Ile Tyr Pro Phe Val Ala Gly Met Thr Gly Ser Ala Leu Ala 405 410 415		
Gly Met Leu Ser Val Thr Phe Asn Val Thr Ala Ala Ser Ile Gly Ile 420 425 430		
Gly Gly Leu Pro Gly Ile Leu Ser Ile Gln Pro Gln Tyr Met Leu Pro 435 440 445		
Phe Ala Gly Thr Met Leu Val Ala Ile Val Val Pro Met Leu Leu Thr 450 455 460		
Phe Phe Phe Arg Lys Ala Gly Leu Phe Thr Lys Thr Glu Gly Asp Thr 465 470 475 480		
Asn Leu Gln Ala Glu Phe Val Ala Gln Glu Glu Ala Glu Phe Val Asn 485 490 495		
His Glu Pro Val Glu Leu Thr Ser Val Glu Ile Ile Ser Pro Leu Thr 500 505 510		
Gly Gln Val Lys Glu Leu Ser Gln Ala Thr Asp Pro Ile Phe Ala Ser 515 520 525		
Gly Val Met Gly Gln Gly Leu Val Ile Glu Pro Ser Gln Gly Glu Leu 530 535 540		

Thr Ser Pro Val Asn Gly Thr Val Thr Val Leu Phe Pro Thr Lys His
 545 550 555 560

Ala Ile Gly Ile Val Ser Asp Glu Gly Val Glu Leu Leu Ile His Ile
 565 570 575

Gly Met Asp Thr Val Gly Leu Asp Gly Lys Gly Phe Glu Ser Leu Val
 580 585 590

Val Gln Gly Asp His Val Thr Val Gly Gln Gln Leu Ile Arg Phe Asp
 595 600 605

Met Asp Val Ile Lys Ala Ala Gly Leu Val Thr Glu Thr Pro Val Ile
 610 615 620

Ile Thr Asn Gln Asp Ala Tyr Thr Ala Thr Ile Pro Gly Thr Tyr Pro
 625 630 635 640

Thr Thr Ile Gln Ala Gly Ala Ser Leu Met Val Ala Thr Arg Ile
 645 650 655

<210> 376

<211> 175

<212> PRT

<213> Streptococcus pneumoniae

<400> 376

Met Ser Arg Val His Val Gln Ile Met Asn Gln Phe His Arg Lys Ser
 1 5 10 15

His Glu Tyr Lys Ala Ile Lys Arg Tyr Trp Lys Leu Ile Gln Gln Asp
 20 25 30

Ser Arg Lys Leu Ser Asp Lys Arg Phe Tyr Arg Pro Thr Phe Arg Met
 35 40 45

His Leu Thr Asn Lys Glu Ile Leu Asp Lys Ile Leu Ser Tyr Ser Glu
 50 55 60

Asp Leu Lys His His Tyr Gln Ile Tyr Gln Leu Leu Leu Phe His Phe
 65 70 75 80

Gln Asn Lys Asp Pro Glu Lys Phe Phe Gly Leu Ile Glu Asp Asn Leu
85 90 95

Lys Gln Val His Pro Leu Phe Gln Thr Val Phe Lys Thr Phe Leu Lys
100 105 110

Asp Lys Glu Lys Ile Ile Asn Ala Leu Gln Leu His Tyr Ser Asn Ala
115 120 125

Lys Leu Glu Ala Thr Asn Asn Leu Ile Lys Leu Ile Lys Arg Asn Ala
130 135 140

Phe Gly Phe Arg Asn Phe Glu Asn Phe Lys Lys Arg Ile Phe Ile Ala
145 150 155 160

Leu Asn Ile Lys Lys Glu Arg Thr Lys Phe Val Leu Ser Arg Ala
165 170 175

<210> 377

<211> 355

<212> PRT

<213> Streptococcus pneumoniae

<400> 377

Met Thr Lys Glu Lys Asn Val Ile Leu Thr Ala Arg Asp Ile Val Val
1 5 10 15

Glu Phe Asp Val Arg Asp Lys Val Leu Thr Ala Ile Arg Gly Val Ser
20 25 30

Leu Glu Leu Val Glu Gly Glu Val Leu Ala Leu Val Gly Glu Ser Gly
35 40 45

Ser Gly Lys Ser Val Leu Thr Lys Thr Phe Thr Gly Met Leu Glu Glu
50 55 60

Asn Gly Arg Ile Ala Gln Gly Ser Ile Asp Tyr Arg Gly Gln Asp Leu
65 70 75 80

Thr Ala Leu Ser Ser His Lys Asp Trp Glu Gln Ile Arg Gly Ala Lys
85 90 95

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Ile Ala Thr   Ile Phe Gln Asp Pro Met Thr Ser Leu Asp Pro Ile Lys
      100                      105                      110

Thr Ile Gly Ser Gln Ile Thr Glu Val Ile Val Lys His Gln Gly Lys
      115                      120                      125

Thr Ala Lys Glu Ala Lys Glu Leu Ala Ile Asp Tyr Met Asn Lys Val
      130                      135                      140

Gly Ile Pro Asp Ala Asp Arg Arg Phe Asn Glu Tyr Pro Phe Gln Tyr
      145                      150                      155                      160

Ser Gly Gly Met Arg Gln Arg Ile Val Ile Ala Ile Ala Leu Ala Cys
      165                      170                      175

Arg Pro Asp Val Leu Ile Cys Asp Glu Pro Thr Thr Ala Leu Asp Val
      180                      185                      190

Thr Ile Gln Ala Gln Ile Ile Asp Leu Leu Lys Ser Leu Gln Asn Glu
      195                      200                      205

Tyr His Phe Thr Thr Ile Phe Ile Thr His Asp Leu Gly Val Val Ala
      210                      215                      220

Ser Ile Ala Asp Lys Val Ala Val Met Tyr Ala Gly Glu Ile Val Glu
      225                      230                      235                      240

Tyr Gly Thr Val Glu Glu Val Phe Tyr Asp Pro Arg His Pro Tyr Thr
      245                      250                      255

Trp Ser Leu Leu Ser Ser Leu Pro Gln Leu Ala Asp Asp Lys Gly Asp
      260                      265                      270

Leu Tyr Ser Ile Pro Gly Thr Pro Pro Ser Leu Tyr Thr Asp Leu Lys
      275                      280                      285

Gly Asp Ala Phe Ala Leu Arg Ser Asp Tyr Ala Met Gln Ile Asp Phe
      290                      295                      300

Glu Gln Lys Ala Pro Gln Phe Ser Val Ser Glu Thr His Trp Ala Lys
      305                      310                      315                      320

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Thr Trp Leu Leu His Glu Asp Ala Pro Lys Val Glu Lys Pro Ala Val
 325 330 335

Ile Ala Asn Leu His Asp Lys Ile Arg Glu Lys Met Gly Phe Ala His
 340 345 350

Leu Ala Asp
 355

<210> 378

<211> 308

<212> PRT

<213> Streptococcus pneumoniae

<400> 378

Met Ser Thr Ile Asp Lys Glu Lys Phe Gln Phe Val Lys Arg Asp Asp
 1 5 10 15

Phe Ala Ser Glu Thr Ile Asp Ala Pro Ala Tyr Ser Tyr Trp Lys Ser
 20 25 30

Val Phe Lys Gln Phe Met Lys Lys Lys Ser Thr Val Val Met Leu Gly
 35 40 45

Ile Leu Val Ala Ile Ile Leu Ile Ser Phe Ile Tyr Pro Met Phe Ser
 50 55 60

Lys Phe Asp Phe Asn Asp Val Ser Lys Val Asn Asp Phe Ser Val Arg
 65 70 75 80

Tyr Ile Lys Pro Asn Ala Glu His Trp Phe Gly Thr Asp Ser Asn Gly
 85 90 95

Lys Ser Leu Phe Asp Gly Val Trp Phe Gly Ala Arg Asn Ser Ile Leu
 100 105 110

Ile Ser Val Ile Ala Thr Val Ile Asn Leu Val Ile Gly Val Phe Val
 115 120 125

Gly Gly Ile Trp Gly Ile Ser Lys Ser Val Asp Arg Val Met Met Glu
 130 135 140

Val Tyr Asn Val Ile Ser Asn Ile Pro Pro Leu Leu Ile Val Ile Val
 145 150 155 160

Leu Thr Tyr Ser Ile Gly Ala Gly Phe Trp Asn Leu Ile Phe Ala Met
 165 170 175

Ser Val Thr Thr Trp Ile Gly Ile Ala Phe Met Ile Arg Val Gln Ile
 180 185 190

Leu Arg Tyr Arg Asp Leu Glu Tyr Asn Leu Ala Ser Arg Thr Leu Gly
 195 200 205

Thr Pro Thr Leu Lys Ile Val Ala Lys Asn Ile Met Pro Gln Leu Val
 210 215 220

Ser Val Ile Val Thr Thr Met Thr Gln Met Leu Pro Ser Phe Ile Ser
 225 230 235 240

Tyr Glu Ala Phe Leu Ser Phe Phe Gly Leu Gly Leu Pro Ile Thr Val
 245 250 255

Pro Ser Leu Gly Arg Leu Ile Ser Asp Tyr Ser Gln Asn Val Thr Thr
 260 265 270

Asn Ala Tyr Leu Phe Trp Ile Pro Leu Thr Thr Leu Val Leu Val Ser
 275 280 285

Leu Ser Leu Phe Val Val Gly Gln Asn Leu Ala Asp Ala Ser Asp Pro
 290 295 300

Arg Thr His Arg
 305

<210> 379
 <211> 498
 <212> PRT
 <213> Streptococcus pneumoniae

<400> 379

Met Lys Lys Tyr Ile Phe Met Arg Val Leu Arg Ser Leu Val Ser Ile
 1 5 10 15

Phe Leu Val Thr Thr Leu Thr Tyr Thr Ile Ile Tyr Thr Leu Val Pro
 20 25 30

Arg Lys Leu Ile Phe Lys Gln Asp Pro Asn Tyr Asn Lys Ile Ala Thr
 35 40 45

Thr Ala Asp Lys Arg Asp Asn Tyr Glu Asn Thr Val Phe Glu Arg Met
 50 55 60

Gly Tyr Ile Glu Tyr Tyr Asp Thr Lys Glu Leu Gln Glu Lys Ala Ser
 65 70 75 80

Ser Met Asp Ser Ser Val Thr Val Glu Ala Asn Ala Thr Asn Lys Ala
 85 90 95

Ile Tyr Glu Lys Tyr Ile Asn Gln Leu Gly His Gly Trp Thr Leu Gly
 100 105 110

Glu Phe Thr Glu Ser Gly Gln Phe Tyr Ala Thr Arg Glu Ile Pro Ile
 115 120 125

Phe Glu Arg Val Phe His Phe Tyr Ala Asn Leu Ile Asp Ile Asp His
 130 135 140

Thr Asn Lys Ile Gln Asp Pro Glu Asn Pro Asp Leu Lys Arg Tyr Leu
 145 150 155 160

Arg Phe Glu Asn Asp Pro Ala Ile Gly Trp Ser Leu Val Gly Ser Gly
 165 170 175

Thr Lys His Lys Tyr Leu Leu Tyr Phe Asn Ser Gln Phe Pro Phe Val
 180 185 190

His Gln Asn Phe Val Asn Leu Asn Leu Gly Asp Ser Tyr Pro Thr Tyr
 195 200 205

Ala Asn Thr Pro Val Leu Gln Val Ile Thr Gln Gly Gln Gly Gln Thr
 210 215 220

Lys Thr Ala Gln Val Gln Phe Pro Thr Gly Lys Lys Thr Ser Ser Val
 225 230 235 240

Asn Ile Tyr Ser Arg Thr Tyr Lys Ser Pro Ser Gln Ala Asp Ser Arg
 245 250 255
 Glu Val Ala Ser Tyr Gly Lys Asp Asp Pro Tyr Thr Ala Thr Glu Ser
 260 265 270
 Asn Tyr Gln Tyr Pro Ser Met Ile Val Ser Ser Ala Ile Thr Gly Leu
 275 280 285
 Ile Gly Leu Val Leu Ala Tyr Ala Leu Ala Val Pro Leu Gly Ser Ala
 290 295 300
 Met Ala Arg Phe Lys Asn Thr Trp Ile Asp Ser Leu Ser Thr Gly Ala
 305 310 315 320
 Leu Thr Phe Leu Leu Ala Leu Pro Thr Ile Ala Leu Val Tyr Ile Val
 325 330 335
 Arg Leu Ile Gly Ser Ser Ile Ala Leu Pro Asp Ser Phe Pro Ile Leu
 340 345 350
 Gly Ala Gly Asp Trp Arg Ser Tyr Val Leu Pro Ala Val Ile Leu Gly
 355 360 365
 Leu Leu Gly Ala Pro Gly Thr Ala Ile Trp Ile Arg Arg Tyr Met Ile
 370 375 380
 Asp Leu Gln Ser Gln Asp Phe Val Arg Phe Ala Arg Ala Lys Gly Leu
 385 390 395 400
 Ser Glu Lys Glu Ile Ser Asn Lys His Ile Phe Lys Asn Ala Met Val
 405 410 415
 Pro Leu Val Ser Gly Ile Pro Ala Ala Ile Ile Gly Val Ile Gly Gly
 420 425 430
 Ala Thr Leu Thr Glu Thr Val Phe Ala Phe Pro Gly Met Gly Lys Met
 435 440 445
 Leu Ile Asp Ser Val Lys Ala Ser Asn Asn Ser Met Val Val Gly Leu

450 455 460
 Val Phe Ile Phe Thr Cys Ile Ser Ile Phe Ser Arg Leu Leu Gly Asp
 465 470 475 480
 Ile Trp Met Thr Ile Ile Asp Pro Arg Ile Lys Leu Thr Glu Lys Gly
 485 490 495
 Gly Lys

 <210> 380
 <211> 343
 <212> PRT
 <213> Streptococcus pneumoniae

 <400> 380
 Met Gly Phe Leu Leu Met Gly Ala Leu Phe Ile Val Leu Pro Arg Thr
 1 5 10 15
 Met Val Ser Ala Lys Arg Ile Asn Gln Val Leu Asp Leu His Ser Ser
 20 25 30
 Ile Gln Asn Pro Val Gln Val Gln Leu Thr Asp Glu Asn Phe Lys Gly
 35 40 45
 Gln Val Glu Phe Lys Asp Val Thr Phe Arg Tyr Ala Ala Asn Ser Glu
 50 55 60
 Ala Val Ile Glu His Val Ser Phe Lys Ala Glu Thr Gly Gln Thr Val
 65 70 75 80
 Ala Phe Ile Gly Ser Thr Gly Ser Gly Lys Ser Thr Leu Val Asn Leu
 85 90 95
 Ile Pro Arg Phe Tyr Asp Val Ser Ala Gly Glu Ile Leu Val Asp Gly
 100 105 110
 Val Asn Val Gln Asp Tyr Asp Phe Ser Ala Thr Ala His Ala Gly Gln
 115 120 125
 Lys Val Ala Ile Val Gly Pro Thr Gly Ala Gly Lys Thr Thr Ile Val

130	135	140	
Asn Leu Leu Met Lys Phe Tyr Glu Ile Asp Lys Gly Ser Ile Arg Ile			
145	150	155	160
Asp Gly Val Asp Thr Lys Ala Met Thr Arg Ser Glu Val His Asp Ala			
	165	170	175
Phe Ser Met Val Leu Gln Asp Thr Trp Leu Phe Glu Gly Thr Ile Arg			
	180	185	190
Asp Asn Leu Ile Tyr Asn Gln Ile Gly Ile Ser Asp Glu Arg Met Met			
	195	200	205
Glu Ala Ser Lys Ala Val Gly Ile His His Phe Ile Met Thr Leu Pro			
	210	215	220
Asp Gly Tyr Asp Thr Ile Leu Asp Asp Thr Val Thr Leu Ser Val Arg			
	225	230	235
Gln Lys Gln Leu Leu Thr Ile Ala Arg Ala Leu Leu Lys Asp Ala Pro			
	245	250	255
Leu Leu Ile Leu Asp Glu Ala Thr Ser Ser Val Asp Thr Arg Thr Glu			
	260	265	270
Glu Leu Ile Gln Lys Ala Met Asp Arg Leu Met Glu Gly Arg Thr Ser			
	275	280	285
Phe Val Ile Ala His Arg Leu Ser Thr Ile Arg Asn Ala Asp Leu Ile			
	290	295	300
Leu Val Met Lys Asp Gly Asn Ile Ile Glu Gln Gly Asn Tyr Glu Glu			
	305	310	315
Leu Met Ala Gln Gly Gly Phe Tyr Ala Asp Leu Tyr Asn Ser Gln Phe			
	325	330	335
Thr Glu Asp Glu Ala Glu Glu			
	340		

<210> 381
 <211> 273
 <212> PRT
 <213> Streptococcus pneumoniae

 <400> 381

 Met Ile Leu Leu Ala Ile Leu Phe Thr Cys Phe Ser Val Tyr Leu Glu
 1 5 10 15

 Leu Glu Val Pro Thr Tyr Ile Ser Lys Ile Thr Asp Leu Leu Gly Ser
 20 25 30

 Gln Glu Thr Asn Leu Asp Glu Leu Trp Gln Ser Ala Ser Met Met Met
 35 40 45

 Gly Met Ser Phe Leu Ala Phe Leu Ser Val Val Ala Val Gly Phe Phe
 50 55 60

 Ala Ser Arg Val Ala Ala Ser Tyr Thr Ser Arg Leu Arg Ser Asp Ile
 65 70 75 80

 Phe Asn Arg Val Leu Asp Tyr Ser Gln Thr Glu Ile Lys Lys Phe Ser
 85 90 95

 Ile Pro Ser Leu Leu Thr Arg Thr Thr Asn Asp Ile Thr Gln Val Gln
 100 105 110

 Met Leu Ile Thr Met Gly Leu Gln Val Val Thr Arg Gly Ser Ile Met
 115 120 125

 Ala Ile Trp Ala Ile Gly Lys Ile Leu Gly His Ser Glu Tyr Trp Leu
 130 135 140

 Trp Ala Val Leu Val Ala Val Ile Ile Asn Val Leu Met Thr Thr Val
 145 150 155 160

 Leu Met Thr Leu Ala Phe Pro Lys Gln Ser Leu Ile Gln Gly Leu Thr
 165 170 175

 Asp Lys Leu Asn Ser Ile Thr Arg Glu Ser Leu Thr Gly Ile Arg Val
 180 185 190

Val Arg Ala Tyr Asn Ala Glu Asp Tyr Gln Asn Glu Lys Phe Ala Ala
 195 200 205

Val Asn Asp Glu Leu Thr Arg Leu Asn Leu Phe Val Asn Arg Leu Met
 210 215 220

Ala Ile Leu Asn Pro Ile Met Met Gly Ile Ser Ser Gly Leu Ser Val
 225 230 235 240

Ala Ile Tyr Trp Ile Gly Ala Tyr Val Ile Asn Asp Ala Ala Pro Ile
 245 250 255

Ala Arg Leu Pro Leu Phe Ser Asp Met Ile Val Phe Met Ser Tyr Ala
 260 265 270

Met

<210> 382

<211> 456

<212> PRT

<213> Streptococcus pneumoniae

<400> 382

Met Asn Lys Lys Arg Thr Val Asp Leu Ile His Gly Pro Ile Leu Pro
 1 5 10 15

Ser Leu Leu Ser Phe Thr Phe Pro Ile Leu Leu Ser Asn Ile Phe Gln
 20 25 30

Gln Leu Tyr Asn Thr Ala Asp Val Leu Ile Val Gly Arg Phe Leu Gly
 35 40 45

Gln Glu Ser Leu Ala Ala Val Gly Ala Thr Thr Ala Ile Phe Asp Leu
 50 55 60

Ile Val Gly Phe Thr Leu Gly Val Gly Asn Gly Met Gly Ile Val Ile
 65 70 75 80

Ala Arg Tyr Tyr Gly Ala Arg Asn Phe Thr Lys Ile Lys Glu Ala Val
 85 90 95

Ala Ala Thr Trp Ile Leu Gly Ala Leu Leu Ser Ile Leu Val Met Leu
 100 105 110

Leu Gly Phe Leu Gly Leu Tyr Pro Leu Leu Gln Tyr Leu Asp Thr Pro
 115 120 125

Ala Glu Ile Leu Pro Gln Ser Tyr Gln Tyr Ile Ser Met Ile Val Thr
 130 135 140

Cys Val Gly Val Ser Phe Ala Tyr Asn Leu Phe Ala Gly Leu Leu Arg
 145 150 155 160

Ser Ile Gly Asp Ser Leu Ala Ala Leu Gly Phe Leu Ile Phe Ser Ala
 165 170 175

Leu Val Asn Val Val Leu Asp Leu Tyr Phe Ile Thr Gln Leu His Leu
 180 185 190

Gly Val Gln Ser Ala Gly Leu Ala Thr Ile Ile Ser Gln Gly Leu Ser
 195 200 205

Ala Val Leu Cys Phe Tyr Tyr Ile Arg Lys Ser Val Pro Glu Leu Leu
 210 215 220

Pro Gln Phe Lys His Phe Lys Trp Asp Lys Ser Leu Tyr Ala Asp Leu
 225 230 235 240

Leu Glu Gln Gly Leu Ala Met Gly Leu Met Ser Ser Ile Val Ser Ile
 245 250 255

Gly Ser Val Ile Leu Gln Phe Ser Val Asn Thr Phe Gly Ala Val Ile
 260 265 270

Ile Ser Ala Gln Thr Ala Ala Arg Arg Ile Met Thr Phe Ala Leu Leu
 275 280 285

Pro Met Thr Ala Ile Ser Ala Ser Met Thr Thr Phe Ala Ser Gln Asn
 290 295 300

Leu Gly Ala Lys Arg Pro Asp Arg Ile Val Gln Gly Leu Arg Ile Gly
 305 310 315 320

Ser Arg Leu Ser Ile Ser Trp Ala Val Phe Val Cys Ile Phe Leu Phe
 325 330 335

Phe Ala Ser Pro Ala Leu Val Ser Phe Leu Ala Ser Ser Thr Asp Gly
 340 345 350

Tyr Leu Ile Glu Asn Gly Ser Leu Tyr Leu Gln Ile Ser Ser Thr Phe
 355 360 365

Tyr Pro Ile Leu Ser Leu Leu Leu Ile Tyr Arg Asn Cys Leu Gln Gly
 370 375 380

Leu Gly Gln Lys Ile Leu Pro Leu Val Ser Ser Phe Ile Glu Leu Ile
 385 390 395 400

Gly Lys Ile Val Phe Val Val Leu Ile Ile Pro Trp Ala Gly Tyr Lys
 405 410 415

Gly Val Ile Leu Cys Glu Pro Leu Ile Trp Val Ala Met Thr Val Gln
 420 425 430

Leu Tyr Phe Ser Leu Phe Arg His Pro Leu Ile Lys Glu Gly Lys Ala
 435 440 445

Ile Leu Ala Thr Lys Val Gln Ser
 450 455

<210> 383

<211> 206

<212> PRT

<213> Streptococcus pneumoniae

<400> 383

Met Lys Ser Met Arg Ile Leu Phe Leu Leu Ala Leu Ile Gln Ile Ser
 1 5 10 15

Leu Ser Ser Cys Phe Leu Trp Lys Glu Cys Ile Leu Ser Phe Lys Gln
 20 25 30

Ser Thr Ala Phe Phe Ile Gly Ser Met Val Phe Val Ser Gly Ile Cys
 35 40 45

Ala Gly Val Asn Tyr Leu Tyr Thr Arg Lys Gln Glu Val His Ser Val
50 55 60

Leu Ala Ser Lys Lys Ser Val Lys Leu Phe Tyr Ser Met Leu Leu Leu
65 70 75 80

Ile Asn Leu Leu Gly Ala Val Leu Val Leu Ser Asp Asn Leu Phe Ile
85 90 95

Lys Asn Thr Leu Gln Gln Glu Leu Val Asp Phe Leu Leu Pro Ser Phe
100 105 110

Phe Phe Leu Phe Gly Leu Asp Leu Leu Ile Phe Leu Pro Leu Lys Lys
115 120 125

Tyr Val Arg Asp Phe Leu Ala Met Leu Asp Arg Lys Lys Thr Val Leu
130 135 140

Val Thr Ile Leu Ala Thr Leu Leu Phe Leu Arg Asn Pro Met Thr Ile
145 150 155 160

Val Ser Leu Leu Ile Tyr Ile Gly Leu Gly Leu Phe Phe Ala Ala Tyr
165 170 175

Leu Val Pro Asn Ser Val Lys Lys Glu Val Ser Phe Tyr Gly His Ile
180 185 190

Phe Arg Asp Leu Val Leu Val Ile Val Thr Leu Ile Phe Phe
195 200 205

<210> 384

<211> 702

<212> PRT

<213> Streptococcus pneumoniae

<400> 384

Met Lys Ile Pro Met Ile Tyr Gln Met Glu Asn Ser Glu Cys Gly Leu
1 5 10 15

Ala Cys Cys Ala Met Ile Leu Asn Tyr Phe Lys Tyr Glu Ile Ser Leu
20 25 30

Asn Glu Leu Arg Glu Ile Tyr Pro Ser Ser Arg Ser Gly Tyr Ser Leu
 35 40 45
 Leu Ser Ile Ser Lys Val Leu Gly Asp Phe Asn Ile Ser Ser His Ala
 50 55 60
 Phe Lys Ala Ser Val Arg Asp Leu Lys Pro Leu Ser Phe Pro Leu Ile
 65 70 75 80
 Cys Phe Trp Glu Ser Ser His Phe Ile Ile Leu Glu Lys Ile Ser Lys
 85 90 95
 Asn Lys Phe Tyr Ile Leu Asp Pro Ala Lys Gly Arg Gln Arg Met Ser
 100 105 110
 Ile Ser Glu Phe Glu Arg His Tyr Ser Asn Ile Ile Leu Thr Phe Lys
 115 120 125
 Lys Leu Asp Ser Phe Met Ser Arg Lys Asp Asn Lys Lys Ser Pro Val
 130 135 140
 Leu Lys Tyr Phe Phe Lys Tyr Arg Asn Lys Leu Gly Ile Leu Phe Phe
 145 150 155 160
 Val Thr Ala Leu Leu Tyr Val Ile Gln Ser Leu Val Pro Ile Ala Asn
 165 170 175
 Arg Tyr Ile Ile Asp Thr Asn Phe Lys Asp Asp Ser Tyr Ser Ser Arg
 180 185 190
 Met Leu Phe Thr Ile Leu Phe Ile Phe Thr Val Ser Phe Ser Leu Met
 195 200 205
 Tyr Leu Leu Arg Gln Ile Tyr Val Ala Ser Leu Lys Tyr Ile Met Asp
 210 215 220
 Lys Glu Ile Ser Tyr Asp Phe Met Lys His Leu Ile Tyr Leu Pro Tyr
 225 230 235 240
 Ser Phe Tyr Glu Lys Arg Thr Leu Gly Asp Ile Leu Phe Arg Ala Asn
 245 250 255

Ser Ile Val Tyr Ile Arg Glu Ile Leu Ser Asn Asn Phe Ile Ala Ala
 260 265 270
 Ile Leu Asp Leu Leu Met Ile Val Val Tyr Ala Val Val Leu Phe Ser
 275 280 285
 Phe Ser Lys Tyr Met Val Ile Phe Leu Ile Ser Leu Ser Leu Ala Leu
 290 295 300
 Ser Ile Val Met Tyr Pro Ile Ile Lys Ile Ser Lys Asn Leu Ile Asp
 305 310 315 320
 Lys Asn Ile Lys Glu Lys Val Asn Val Gln Asn Ile Thr Ser Glu Val
 325 330 335
 Ile Ser Lys Asn Ser Asp Ile Lys Leu Thr Gly Glu Glu Glu Phe Trp
 340 345 350
 Ile Asn Lys Trp Asp Asn Phe Asn Thr Lys Gln Leu Ile Ile Gly Arg
 355 360 365
 Lys Leu Asp Ile His Leu Ser Ile Val Ser Ser Ile Thr Asn Val Leu
 370 375 380
 Gln Ile Ile Leu Pro Val Leu Thr Leu Ile Val Gly Val Asn Ile Lys
 385 390 395 400
 Thr Phe Glu Gln Leu Thr Leu Gly Gln Ile Val Ala Ile Ser Thr Val
 405 410 415
 Ser Pro Tyr Phe Ile Ser Pro Ile Ile Ser Leu Ser Asp Asn Tyr Ile
 420 425 430
 Gln Leu Met Leu Leu Lys Gly Tyr Phe Leu Arg Ile Glu Asp Val Phe
 435 440 445
 Asn Thr Lys Ser Glu Leu Ile Pro Glu Arg Val Ser Gln Asp Ile Lys
 450 455 460
 Phe Asp Lys Lys Ile Glu Leu Lys Asp Ile Trp Tyr Lys Tyr Gly Leu

465	470	475	480
Phe Asp Asp Tyr Val Leu Lys Gly Ile Asn Val Thr Ile Lys Lys Gly	485	490	495
Glu Thr Val Ala Ile Val Gly Glu Ser Gly Ser Gly Lys Ser Thr Leu	500	505	510
Ala Lys Ile Leu Leu Gly Leu Leu Glu Pro Asn Ile Gly Ser Ile Glu	515	520	525
Val Asp Gly Val Glu Lys Glu Glu Ile Gly Gln Thr Leu Tyr Arg Lys	530	535	540
Ile Phe Gly Ala Val Leu Gln Asn Ser Thr Leu Ser Tyr Gly Thr Leu	545	550	555
Arg Glu Asn Leu Thr Phe Gly His Phe Val Ser Asp Glu Glu Leu Met	565	570	575
Thr Asn Leu Asn Ser Ile Gly Leu Ser Asn Val Val Lys Ser Leu Pro	580	585	590
Leu Gly Leu Glu Thr Ile Ile Ala Glu Glu Gly Asn Asn Phe Ser Gly	595	600	605
Gly Gln Gln Gln Met Ile Leu Leu Ala Arg Cys Leu Leu Ser Lys Pro	610	615	620
Ser Val Val Val Leu Asp Glu Ala Thr Ser Ser Leu Asp Asn Leu Ser	625	630	635
Gln Gln Ile Thr Thr Ser Tyr Leu Ser Glu Ile Gly Thr Thr Lys Ile	645	650	655
Leu Ile Ala His Arg Leu Asp Thr Ile Lys Ser Ala Asp Lys Ile Leu	660	665	670
Val Met His Asn Gly Glu Ile Val Glu Ile Gly Thr His Arg Glu Leu	675	680	685

Leu Glu Leu Gly Gly Ile Tyr Lys Gln Leu Tyr Ser Asn Asn
 690 695 700

<210> 385
 <211> 103
 <212> PRT
 <213> Streptococcus pneumoniae

<400> 385

Met Leu Asn Ile Pro Asn Val Leu Arg Tyr Asp Leu Asn Met Leu Gln
 1 5 10 15

Leu Glu Tyr Lys Asn Glu Gln Ser Trp Asp Ser Phe Ile Asp Asn Val
 20 25 30

Asn Leu Ile Glu Leu Glu Glu Arg Ile Gln Thr Thr Ile Gly Ile Lys
 35 40 45

Gln Ile Asn Thr His Asn Ile Ile Thr Ile Ala Arg Glu Gly Tyr Ser
 50 55 60

Gln Asn Tyr Leu Pro Asn Thr Ser Glu Asn Thr Tyr Asn Ser Leu Gln
 65 70 75 80

Val Ser Leu Val Gly Val Leu Leu Leu Phe Ile Ser Met Val Asn Ile
 85 90 95

Leu Trp Ala Lys Lys Ser Lys
 100

<210> 386
 <211> 210
 <212> PRT
 <213> Streptococcus pneumoniae

<400> 386

Met Ile Glu Leu Lys Gln Val Ser Lys Ser Phe Gly Glu Arg Glu Leu
 1 5 10 15

Phe Ser Asn Leu Ser Met Thr Phe Glu Ala Gly Lys Val Tyr Ala Leu
 20 25 30

Ile Gly Ser Ser Gly Ser Gly Lys Thr Thr Leu Met Asn Met Ile Gly

35	40	45
Lys Leu Glu Pro Tyr Asp 50	Gly Thr Ile Phe Tyr Arg 55	Gly Lys Asp Leu 60
Ala Asn Tyr Lys Ser Ser Asp Phe Phe Arg His Glu Leu Gly Tyr Leu 65	70	75 80
Phe Gln Asn Phe Gly Leu Ile Glu Asn Gln Ser Ile Glu Glu Asn Leu 85	90	95
Lys Leu Gly Leu Ile Gly Gln Lys Leu Ser Arg Ser Glu Gln Arg Leu 100	105	110
Arg Gln Lys Gln Ala Leu Glu Gln Val Gly Leu Val Tyr Leu Asp Leu 115	120	125
Asp Lys Arg Ile Phe Glu Leu Ser Gly Gly Glu Ser Gln Arg Val Ala 130	135	140
Leu Ala Lys Ile Ile Leu Lys Asn Pro Pro Phe Ile Leu Ala Asp Glu 145	150	155 160
Pro Thr Ala Ser Ile Asp Pro Ala Thr Ser Gln Leu Ile Met Glu Ile 165	170	175
Leu Leu Ser Leu Arg Asp Asp Asn Arg Leu Ile Ile Ile Ala Thr His 180	185	190
Asn Pro Ala Ile Trp Glu Met Ala Asp Glu Val Phe Thr Met Asp His 195	200	205
Leu Lys 210		
<210> 387		
<211> 345		
<212> PRT		
<213> Streptococcus pneumoniae		
<400> 387		
Met Lys Lys Lys Ile Arg Trp Pro Leu Tyr Val Ile Ala Ala Leu Ile		

1	5	10	15
Val Thr Phe	Leu Ala Phe	Val Val Pro	Leu Pro Tyr Tyr Ile Glu Val
	20	25	30
Pro Gly Gly	Ser Glu Asp	Ile Arg Gln	Val Leu Lys Val Asn Asp Thr
	35	40	45
Glu Asp Lys	Glu Ala Gly	Ala Tyr Gln	Phe Val Thr Val Gly Val Gln
	50	55	60
His Ala Thr	Leu Ala His	Met Ile Tyr	Ala Trp Leu Thr Pro Phe Thr
	65	70	75
Asp Ile Arg	Ser Ala Gln	Glu Thr Thr	Gly Gly Ser Ser Asp Val Glu
	85	90	95
Phe Met Arg	Ile Asn Gln	Phe Tyr Met	Gln Thr Ser Gln Asn Met Ala
	100	105	110
Lys Tyr Gln	Gly Leu Lys	Thr Ala Gly	Lys Asp Ile Glu Leu Lys Tyr
	115	120	125
Phe Gly Val	Tyr Val Leu	Asn Val Thr	Asp Asn Ser Thr Phe Lys Gly
	130	135	140
Ile Leu Asn	Ile Ser Asp	Thr Val Thr	Ala Val Asn Asp Gln Thr Phe
	145	150	155
Asp Ser Ser	Lys Asp Leu	Ile Asp Tyr	Val Ser Ser Gln Lys Leu Gly
	165	170	175
Asp Ser Val	Lys Val Thr	Tyr Glu Glu	Asp Gly Gln Thr Lys Ser Ala
	180	185	190
Glu Gly Lys	Ile Ile Thr	Leu Glu Asn	Gly Lys Asn Gly Ile Gly Ile
	195	200	205
Gly Leu Ile	Asp Arg Thr	Glu Val Ile	Ser Asn Val Pro Ile Ser Phe
	210	215	220

Ser Thr Ala Gly Ile Gly Gly Pro Ser Ala Gly Leu Met Phe Ser Leu
 225 230 235 240

Ala Ile Tyr Thr Gln Ile Ala His Pro Asp Leu Arg Asn Gly Arg Ile
 245 250 255

Val Ala Gly Thr Gly Thr Ile Asp Arg Asp Gly Asn Val Gly Asp Ile
 260 265 270

Gly Gly Ile Asp Lys Lys Val Val Ala Ser Ala Arg Ala Gly Ala Ala
 275 280 285

Ile Phe Phe Ala Pro Asp Asn Pro Val Ser Glu Glu Glu Gln Lys Ala
 290 295 300

His Pro Asp Ala Lys Asn Asn Tyr Gln Thr Ala Leu Glu Ala Ala Lys
 305 310 315 320

Thr Ile Lys Thr Asp Met Lys Ile Val Pro Val Lys Thr Leu Gln Asp
 325 330 335

Ala Ile Asp Tyr Leu Lys Asn Asn Pro
 340 345

<210> 388
 <211> 308
 <212> PRT
 <213> Streptococcus pneumoniae

<400> 388

Met Lys Ser Ile Lys Arg Phe Ala Leu Ser Ala Met Gly Val Ala Met
 1 5 10 15

Leu Leu Val Leu Thr Gly Cys Val Asn Val Asp Lys Thr Thr Gly Gln
 20 25 30

Pro Thr Gly Phe Ile Trp Asn Thr Ile Gly Ala Pro Met Ala Glu Ala
 35 40 45

Ile Lys Tyr Phe Ala Thr Asp Lys Gly Leu Gly Phe Gly Val Ala Ile
 50 55 60

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Ile Ile Val Thr Ile Ile Val Arg Leu Ile Ile Leu Pro Leu Gly Ile
65              70              75              80

Tyr Gln Ser Trp Lys Ala Thr Leu His Ser Glu Lys Met Asn Ala Leu
85              90              95

Lys His Val Leu Glu Pro His Gln Thr Arg Leu Lys Glu Ala Thr Thr
100             105             110

Gln Glu Glu Lys Leu Glu Ala Gln Gln Ala Leu Phe Ala Ala Gln Lys
115             120             125

Glu His Gly Ile Ser Met Phe Gly Gly Val Gly Cys Phe Pro Ile Leu
130             135             140

Leu Gln Met Pro Phe Phe Ser Ala Ile Tyr Phe Ala Ala Gln His Thr
145             150             155             160

Glu Gly Val Ala Gln Ala Ser Tyr Leu Gly Ile Pro Leu Gly Ser Pro
165             170             175

Ser Met Ile Leu Val Ala Cys Ala Gly Val Leu Tyr Tyr Leu Gln Ser
180             185             190

Leu Leu Ser Leu His Gly Val Glu Asp Glu Met Gln Arg Glu Gln Ile
195             200             205

Lys Lys Met Ile Tyr Met Ser Pro Leu Met Ile Val Val Phe Ser Leu
210             215             220

Phe Ser Pro Ala Ser Val Thr Leu Tyr Trp Val Val Gly Gly Phe Met
225             230             235             240

Met Ile Leu Gln Gln Phe Ile Val Asn Tyr Ile Val Arg Pro Lys Leu
245             250             255

Arg Lys Lys Val Arg Glu Glu Leu Ala Lys Asn Pro Pro Lys Ala Ser
260             265             270

Ala Phe Ser Lys Pro Ser Gly Arg Lys Asp Val Thr Pro Glu Gln Pro
275             280             285

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Thr Ala Ile Thr Ser Lys Lys Lys His Lys Asn Arg Asn Ala Gly Lys
 290 295 300

Gln Arg Ser Arg
 305

<210> 389
 <211> 213
 <212> PRT
 <213> Streptococcus pneumoniae

<400> 389

Met Ile Asp Ile Gln Gly Leu Glu Lys Lys Phe Asn Asp Arg Ala Ile
 1 5 10 15

Phe Ser Gly Leu Asn Leu Lys Leu Glu Lys Gly Lys Val Tyr Ala Leu
 20 25 30

Ile Gly Lys Ser Gly Ser Gly Lys Thr Thr Leu Leu Asn Ile Leu Gly
 35 40 45

Lys Leu Glu Lys Ile Asp Gly Gly Arg Val Leu Tyr Gln Gly Lys Asp
 50 55 60

Leu Lys Thr Ile Pro Thr Arg Glu Tyr Phe Arg Asp Gln Met Gly Tyr
 65 70 75 80

Leu Phe Gln Asn Phe Gly Leu Leu Glu Asn Gln Ser Ile Lys Glu Asn
 85 90 95

Leu Asp Leu Gly Phe Val Gly Gln Lys Ile Ser Lys Val Glu Arg Leu
 100 105 110

Glu Arg Gln Val Gly Ala Leu Glu Lys Val Asn Leu Gly Tyr Leu Asp
 115 120 125

Leu Glu Gln Lys Ile Tyr Thr Leu Ser Gly Gly Glu Ala Gln Arg Val
 130 135 140

Ala Leu Ala Lys Thr Ile Leu Lys Asn Pro Pro Leu Ile Leu Ala Asp
 145 150 155 160

Glu Pro Thr Ala Ala Leu Asp Pro Glu Asn Ser Glu Glu Val Met Asn
 165 170 175

Leu Leu Val Asp Leu Lys Asp Glu Asn Arg Ile Ile Ile Ile Ala Thr
 180 185 190

His Asn Pro Leu Val Trp Asn Lys Ala Asp Glu Ile Ile Asp Met Arg
 195 200 205

Lys Leu Ala His Val
 210

<210> 390
 <211> 680
 <212> PRT
 <213> Streptococcus pneumoniae

<400> 390

Met Glu Glu Met Met Lys Arg Leu Phe Ile Leu Ile Ser Met Val Leu
 1 5 10 15

Val Ser Leu Tyr Met Val Ile Thr Ser Val Asp His Arg Glu Glu Ile
 20 25 30

Leu Phe Gly Asn Tyr Pro Ser Val Asp Val Thr Gly Met Met Ile Asn
 35 40 45

Gln Pro Val Ala Ser Arg Glu Glu Val Thr Glu Ala Leu Ser His Leu
 50 55 60

Ala Val Glu His Asn Ser Leu Ile Ala Arg Arg Ile Val Glu Pro Asn
 65 70 75 80

Glu Ala Gly Glu Thr Arg Phe Thr Tyr Ala Thr Tyr Gly Glu Gly Lys
 85 90 95

Leu Pro Glu Gly Leu Thr Ile Ser Ser Lys Glu Ser Ala Glu Thr Ser
 100 105 110

Asp Leu Leu Gly Ser Tyr Leu Ile Val Ser Gly Ser Leu Asp Gly Val
 115 120 125

Ser Leu Gln Thr Thr Leu Lys Glu Leu Gly Tyr Gln Gly Phe Val Ser
 130 135 140
 Asn Gly Glu Asp Pro Phe Ser Ile Val Leu Leu Leu Thr Ala Thr Pro
 145 150 155 160
 Met Val Leu Leu Ser Leu Ala Ile Phe Leu Leu Thr Phe Met Ser Leu
 165 170 175
 Thr Leu Ile Tyr Arg Ile Lys Ser Leu Arg Gln Ala Gly Ile Arg Leu
 180 185 190
 Ile Ala Gly Glu Ser Leu Phe Gly Val Ala Leu Arg Pro Val Leu Glu
 195 200 205
 Asp Val Arg Gln Leu Ile Cys Ser Val Leu Val Ser Ser Leu Leu Gly
 210 215 220
 Leu Gly Ile Leu Trp Tyr Gln Gly Ala Leu Phe Met Ala Thr Val Gln
 225 230 235 240
 Leu Val Ile Ile Ala Leu Leu Leu Tyr Gly Leu Thr Leu Ala Gly Ile
 245 250 255
 Ser Thr Leu Leu Ser Val Val Tyr Leu Leu Gly Leu Gln Glu Asn Ser
 260 265 270
 Leu Val Asp Leu Leu Lys Gly Lys Leu Pro Leu Lys Arg Met Met Thr
 275 280 285
 Leu Met Met Val Gly Gln Leu Leu Ala Val Leu Val Val Gly Ser Ser
 290 295 300
 Ala Thr Ala Leu Leu Pro His Tyr Arg Glu Met Gln Glu Met Glu Arg
 305 310 315 320
 Ala Ser Asn Lys Trp Ser Gln Ser Ser Asp Arg Tyr Arg Leu Ser Phe
 325 330 335
 Gly Trp Ser Ser Ala Phe Ala Asp Glu Glu Gly Thr Arg Lys Asp Asn
 340 345 350

Arg Glu Trp Gln Thr Phe Thr Glu Glu Arg Leu Ala Asn Thr Asp Ser
 355 360 365
 Phe Tyr Ile Met Ser Asn Val Asp Asn Phe Ser Asp Gly Ala Glu Val
 370 375 380
 Asp Leu Asp Gly Asn Arg Leu Ser Asp Tyr Thr Pro Ser Gly Asn Val
 385 390 395 400
 Ile Tyr Val Ser Pro Arg Tyr Leu Ile Glu Glu Lys Ile Thr Val Ser
 405 410 415
 Ser Glu Phe Met Asp Lys Met Gln Asn Leu Ser Glu Gly Glu Phe Gly
 420 425 430
 Leu Ile Leu Pro Glu Ser Leu Arg Glu Gln Ser Val Tyr Tyr Gln Gly
 435 440 445
 Leu Phe Thr Asp Tyr Leu Gln Asn Phe Ser Ser Glu Ser Val Glu Val
 450 455 460
 Thr Ser Gln Lys His Tyr Leu Pro Gln Val Arg Leu Ala Phe Thr Glu
 465 470 475 480
 Thr Gly Gln Glu Arg Phe Leu Tyr Asn Asp Gly Tyr Lys Thr Thr Arg
 485 490 495
 Gln Tyr Leu Lys Asp Pro Ile Ile Val Val Leu Thr Pro Gln Ala Thr
 500 505 510
 Gly Thr Arg Pro Val Ala Gly Met Leu Trp Gly Thr Thr Ala Asn Ser
 515 520 525
 Ala Leu Lys Leu Asp Arg Tyr Gly Asp Ser Ile Thr Ala Leu Lys Glu
 530 535 540
 Lys Gly Leu Tyr His Lys Val Ser Tyr Leu Val Lys Ser Gln Leu Phe
 545 550 555 560
 Phe Ala Lys Val Leu Asn Asp Lys Arg Val Glu Phe Tyr Ser Leu Leu

565 570 575
 Ile Gly Thr Ile Leu Thr Leu Ser Thr Ala Ile Leu Leu Phe Asp Ser
 580 585 590
 Met Asn Leu Leu Tyr Phe Glu Gln Phe Arg Arg Glu Leu Met Ile Lys
 595 600 605
 Arg Leu Ala Gly Met Thr Ile Tyr Glu Leu His Gly Lys Tyr Leu Leu
 610 615 620
 Ala Gln Gly Gly Val Leu Leu Leu Gly Leu Val Leu Ser Ser Ile Leu
 625 630 635 640
 Thr Arg Asp Gly Leu Ile Ser Ala Leu Val Val Ala Leu Phe Thr Leu
 645 650 655
 Asn Ala Leu Leu Ile Leu Val Arg Gln Asp Lys Lys Glu Glu Ala Gly
 660 665 670
 Ser Met Ala Val Leu Lys Gly Lys
 675 680
 <210> 391
 <211> 294
 <212> PRT
 <213> Streptococcus pneumoniae
 <400> 391
 Met Asn Met Ile Lys Val Glu Ser Leu Asn Lys Asn Ile Lys Gly Lys
 1 5 10 15
 Ala Ile Leu Lys Gly Ile Ser Phe Glu Val Ala Glu Gly Glu Cys Val
 20 25 30
 Ala Leu Ile Gly Pro Asn Gly Ala Gly Lys Thr Thr Leu Leu Asp Cys
 35 40 45
 Leu Leu Gly Asp Lys Leu Val Thr Ser Gly Gln Val Ser Ile Gln Gly
 50 55 60
 Leu Ser Val Thr Ser Ser Gln Leu Asp Tyr Ile Arg Gly Tyr Leu Pro

65		70		75		80
Gln Glu Asn Val	Ile Val Gln Lys	Leu Lys Val Lys	Glu Leu Ile Ala			
	85		90			95
Phe Phe Gln Arg	Ile Tyr Pro Asn Ser	Leu Ser Asp Gln	Glu Ile Asp			
	100		105			110
Gln Leu Leu Gln	Phe Asp Gln Gln	Gln Lys Glu Gln	Phe Ala Glu Lys			
	115		120			125
Leu Ser Gly Gly	Gln Lys Arg Leu	Phe Ser Phe Val	Leu Thr Leu Ile			
	130		135			140
Gly Arg Pro Lys	Leu Val Phe Leu	Asp Glu Pro Thr	Ala Ala Met Asp			
	145		150			155
Thr Ser Thr Arg	Gln Arg Phe Trp	Glu Ile Val Arg	Asp Leu Lys Ala			
	165		170			175
Gln Gly Val Thr	Ile Leu Tyr Ser	Ser His Tyr Ile	Glu Glu Val Glu			
	180		185			190
His Thr Ala Asp	Arg Ile Leu Val	Leu Asn Lys Gly	Glu Leu Ile Arg			
	195		200			205
Asp Thr Thr Pro	Leu Ala Met Arg	Ser Glu Gly Ile	Glu Lys His Phe			
	210		215			220
Ile Leu Pro Leu	Ala Tyr Lys Glu	Val Ile Glu Gln	Ser Asn Leu Val			
	225		230			235
Glu Asn Trp Ser	Gln Lys Gln Asp	Ala Leu Gln Val	Val Thr Arg Glu			
	245		250			255
Ala Asp Ala Phe	Trp Glu Leu Leu	Val Gln Ala Gly	Cys Gly Ile Gln			
	260		265			270
Glu Ile Glu Val	Asn Asn Arg Ser	Leu Leu Asp Thr	Ile Phe Glu Glu			
	275		280			285

Thr Gln Lys Gly Asp Asn
290

<210> 392

<211> 431

<212> PRT

<213> Streptococcus pneumoniae

<400> 392

Met Lys Asn Ser Lys Phe Ile Asp Gln Phe Ala Thr Phe Ala Gly Lys
1 5 10 15

Leu Gly Asn Gln Ile His Leu Lys Thr Leu Arg Asp Ala Phe Val Thr
20 25 30

Val Met Pro Leu Tyr Ile Leu Ala Gly Leu Ile Val Leu Leu Asn Asn
35 40 45

Thr Val Phe Lys Trp Ile Phe Gln Gly Asp Thr Leu Thr Arg Phe Gln
50 55 60

Tyr Trp Gly Ile Thr Ile Ala Asn Gly Thr Leu Ser Ile Ser Gly Met
65 70 75 80

Ile Ile Ala Val Met Val Gly Tyr Phe Leu Ala Lys Asn Arg Asp Phe
85 90 95

Glu Asn Pro Leu Ala Ala Ser Met Leu Ser Leu Val Ser Leu Ile Val
100 105 110

Met Met Pro Asn Thr Val Ser Val Val Pro Asp Gly Ala Lys Asp Ala
115 120 125

Val Asn Ile Ser Gly Val Leu Ser Phe Asn Asn Thr Gly Thr Gly Ala
130 135 140

Met Phe Ala Gly Val Ile Val Ala Ile Ile Ala Thr Glu Leu Phe Ile
145 150 155 160

Glu Leu Ser Asn Val Lys Ala Leu Gln Met Asn Leu Gly Glu Asn Ile
165 170 175

Pro Pro Ala Val Ser Arg Ser Phe Ser Val Leu Leu Pro Val Met Thr
 180 185 190

Val Ile Ser Leu Phe Gly Val Val Ser Ala Leu Leu Phe Asn Ile Thr
 195 200 205

Gly Met Asn Leu Ile Ser Ile Ile Thr Ile Phe Ile Gln Glu Pro Ile
 210 215 220

Arg His Ile Gly Thr Ser Leu Ile Gly Val Ile Ile Ile Tyr Ser Leu
 225 230 235

Gly Asn Met Leu Trp Leu Phe Gly Ile His Gln Ala Val Ile Tyr Ser
 245 250 255

Ala Ile Leu Glu Pro Leu Leu Leu Ile Asn Ile Thr Glu Asn Ile Thr
 260 265 270

Ala Ala Asn Asn Gly Gln Ala Ile Pro His Ile Ile Asn Leu Ser Gln
 275 280 285

Ile Gln Thr Phe Ala Leu Met Gly Gly Ser Gly Ser Thr Leu Cys Leu
 290 295 300

Leu Ile Ala Thr Phe Leu Val Ser Arg Asn Ala Val Ser Lys Asn Val
 305 310 315 320

Ala Lys Leu Ser Phe Gly Pro Gly Ile Phe Asn Ile Asn Glu Pro Val
 325 330 335

Leu Phe Gly Tyr Pro Ile Val Tyr Asn Ile Ser Leu Ala Ile Pro Phe
 340 345 350

Ile Thr Val Pro Val Leu Gly Ile Leu Ile Ser Tyr Leu Ala Thr Val
 355 360 365

Thr Glu Phe Met Ser Pro Ala Phe Ile Gln Val Pro Trp Thr Thr Pro
 370 375 380

Val Phe Leu Asn Ala Trp Leu Ala Thr Ala Gly Asp Val Arg Ala Val
 385 390 395 400

Leu Val Gln Phe Ile Ile Phe Ala Leu Gly Val Leu Leu Tyr Ile Pro
 405 410 415

Phe Ile Lys Val Asn Asp Lys Val Val Glu Gln Glu Met Glu Gly
 420 425 430

<210> 393

<211> 890

<212> PRT

<213> Streptococcus pneumoniae

<400> 393

Met Lys Ala Met Glu Glu Asn Met Ala Asp Lys Lys Thr Val Thr Pro
 1 5 10 15

Glu Glu Lys Lys Leu Val Ala Glu Lys His Val Asp Glu Leu Val Gln
 20 25 30

Lys Ala Leu Val Ala Leu Glu Glu Met Arg Lys Leu Asp Gln Glu Gln
 35 40 45

Val Asp Tyr Ile Val Ala Lys Ala Ser Val Ala Ala Leu Asp Ala His
 50 55 60

Gly Glu Leu Ala Leu His Ala Phe Glu Glu Thr Gly Arg Gly Val Phe
 65 70 75 80

Glu Asp Lys Ala Thr Lys Asn Leu Phe Ala Cys Glu His Val Val Asn
 85 90 95

Asn Met Arg His Thr Lys Thr Val Gly Val Ile Glu Glu Asp Asp Val
 100 105 110

Thr Gly Leu Thr Leu Ile Ala Glu Pro Val Gly Val Val Cys Gly Ile
 115 120 125

Thr Pro Thr Thr Asn Pro Thr Ser Thr Ala Ile Phe Lys Ser Leu Ile
 130 135 140

Ser Leu Lys Thr Arg Asn Pro Ile Val Phe Ala Phe His Pro Ser Ala
 145 150 155 160

Gln Glu Ser Ser Ala His Ala Ala Arg Ile Val Arg Asp Ala Ala Ile
 165 170 175
 Ala Ala Gly Ala Pro Glu Asn Cys Val Gln Trp Ile Thr Gln Pro Ser
 180 185 190
 Met Glu Ala Thr Ser Ala Leu Met Asn His Glu Gly Val Ala Thr Ile
 195 200 205
 Leu Ala Thr Gly Gly Asn Ala Met Val Lys Ala Ala Tyr Ser Cys Gly
 210 215 220
 Lys Pro Ala Leu Gly Val Gly Ala Gly Asn Val Pro Ala Tyr Val Glu
 225 230 235 240
 Lys Ser Ala Asn Ile Arg Gln Ala Ala His Asp Ile Val Met Ser Lys
 245 250 255
 Ser Phe Asp Asn Gly Met Val Cys Ala Ser Glu Gln Ala Val Ile Ile
 260 265 270
 Asp Lys Glu Ile Tyr Asp Glu Phe Val Ala Glu Phe Lys Ser Tyr His
 275 280 285
 Thr Tyr Phe Val Asn Lys Lys Glu Lys Ala Leu Leu Glu Glu Phe Cys
 290 295 300
 Phe Gly Val Lys Ala Asn Ser Lys Asn Cys Ala Gly Ala Lys Leu Asn
 305 310 315 320
 Ala Asp Ile Val Gly Lys Pro Ala Thr Trp Ile Ala Glu Gln Ala Gly
 325 330 335
 Phe Thr Val Pro Glu Gly Thr Asn Ile Leu Ala Ala Glu Cys Lys Glu
 340 345 350
 Val Gly Glu Asn Glu Pro Leu Thr Arg Glu Lys Leu Ser Pro Val Ile
 355 360 365
 Ala Val Leu Lys Ser Glu Ser Arg Glu Asp Gly Ile Thr Lys Ala Arg
 370 375 380

Gln Met Val Glu Phe Asn Gly Leu Gly His Ser Ala Ala Ile His Thr
 385 390 395 400
 Ala Asp Glu Glu Leu Thr Lys Glu Phe Gly Lys Ala Val Lys Ala Ile
 405 410 415
 Arg Val Ile Cys Asn Ser Pro Ser Thr Phe Gly Gly Ile Gly Asp Val
 420 425 430
 Tyr Asn Ala Phe Leu Pro Ser Leu Thr Leu Gly Cys Gly Ser Tyr Gly
 435 440 445
 Arg Asn Ser Val Gly Asp Asn Val Ser Ala Ile Asn Leu Leu Asn Ile
 450 455 460
 Lys Lys Val Gly Arg Arg Arg Asn Asn Met Gln Trp Met Lys Leu Pro
 465 470 475 480
 Ser Lys Thr Tyr Phe Glu Arg Asp Ser Ile Gln Tyr Leu Gln Lys Cys
 485 490 495
 Arg Asp Val Glu Arg Val Met Ile Val Thr Asp His Ala Met Val Glu
 500 505 510
 Leu Gly Phe Leu Asp Arg Ile Ile Glu Gln Leu Asp Leu Arg Arg Asn
 515 520 525
 Lys Val Val Tyr Gln Ile Phe Ala Asp Val Glu Pro Asp Pro Asp Ile
 530 535 540
 Thr Thr Val Asn Arg Gly Thr Glu Ile Met Arg Ala Phe Lys Pro Asp
 545 550 555 560
 Thr Ile Ile Ala Leu Gly Gly Gly Ser Pro Met Asp Ala Ala Lys Val
 565 570 575
 Met Trp Leu Phe Tyr Glu Gln Pro Glu Val Asp Phe Arg Asp Leu Val
 580 585 590
 Gln Lys Phe Met Asp Ile Arg Lys Arg Ala Phe Lys Phe Pro Leu Leu

595	600	605
Gly Lys Lys Thr Lys Phe Ile Ala Ile Pro Thr Thr Ser Gly Thr Gly 610 615 620		
Ser Glu Val Thr Pro Phe Ala Val Ile Ser Asp Lys Ala Asn Asn Arg 625 630 635 640		
Lys Tyr Pro Ile Ala Asp Tyr Ser Leu Thr Pro Thr Val Ala Ile Val 645 650 655		
Asp Pro Ala Leu Val Leu Thr Val Pro Gly Phe Val Ala Ala Asp Thr 660 665 670		
Gly Met Asp Val Leu Thr His Ala Thr Glu Ala Tyr Val Ser Gln Met 675 680 685		
Ala Ser Asp Tyr Thr Asp Gly Leu Ala Leu Gln Ala Ile Lys Leu Val 690 695 700		
Phe Glu Asn Leu Glu Ser Ser Val Lys Asn Ala Asp Phe His Ser Arg 705 710 715 720		
Glu Lys Met His Asn Ala Ser Thr Ile Ala Gly Met Ala Phe Ala Asn 725 730 735		
Ala Phe Leu Gly Ile Ser His Ser Met Ala His Lys Ile Gly Ala Gln 740 745 750		
Phe His Thr Ile His Gly Arg Thr Asn Ala Ile Leu Leu Pro Tyr Val 755 760 765		
Ile Arg Tyr Asn Gly Thr Arg Pro Ala Lys Thr Ala Thr Trp Pro Lys 770 775 780		
Tyr Asn Tyr Tyr Arg Ala Asp Glu Lys Tyr Gln Asp Ile Ala Arg Met 785 790 795 800		
Leu Gly Leu Pro Ala Ser Thr Pro Glu Glu Gly Val Glu Ser Tyr Ala 805 810 815		

Lys Ala Val Tyr Glu Leu Gly Glu Arg Ile Gly Ile Gln Met Asn Phe
820 825 830

Arg Asp Gln Gly Ile Asp Glu Lys Glu Trp Lys Glu His Ser Arg Lys
835 840 845

Leu Ala Phe Leu Ala Tyr Glu Asp Gln Cys Ser Pro Ala Asn Pro Arg
850 855 860

Leu Pro Met Val Asp His Met Gln Glu Ile Ile Glu Asp Ala Tyr Tyr
865 870 875 880

Gly Tyr Lys Glu Arg Pro Gly Arg Arg Lys
885 890

<210> 394

<211> 99

<212> PRT

<213> Streptococcus pneumoniae

<400> 394

Met Asn Pro Asn Ile Thr Phe Leu Ile Met Leu Val Gly Met Met Ala
1 5 10 15

Leu Met Phe Phe Met Gln Arg Ser Gln Lys Lys Gln Ala Gln Lys Arg
20 25 30

Met Glu Ser Leu Asn Lys Leu Gln Lys Gly Tyr Glu Val Ile Thr Ile
35 40 45

Gly Gly Leu Tyr Gly Thr Val Asp Glu Val Asp Thr Glu Lys Gly Thr
50 55 60

Ile Val Leu Asp Val Asp Gly Val Tyr Leu Thr Phe Glu Leu Ala Ala
65 70 75 80

Ile Lys Thr Val Leu Pro Leu Lys Glu Thr Ala Ser Leu Glu Gly Ala
85 90 95

Ile Glu Lys

<210> 395

<211> 605

<212> PRT

<213> Streptococcus pneumoniae

<400> 395

Met Arg Ile Lys Trp Phe Ser Leu Ile Arg Ile Ile Gly Leu Leu Leu
 1 5 10 15

Val Leu Leu Tyr His Phe Phe Gln Thr Ile Phe Pro Gly Gly Phe Phe
 20 25 30

Gly Val Asp Val Phe Phe Thr Phe Ser Gly Phe Leu Ile Thr Ala Leu
 35 40 45

Leu Ile Glu Glu Phe Ser Lys Asn Asn Glu Ile Asp Leu Ile Gly Phe
 50 55 60

Phe Arg Arg Arg Phe Tyr Arg Ile Val Pro Pro Val Val Leu Met Val
 65 70 75 80

Leu Val Thr Met Pro Phe Thr Phe Leu Val Arg Gln Asp Tyr Val Ala
 85 90 95

Gly Ile Gly Gly Gln Ile Ala Gly Val Leu Gly Phe Met Thr Asn Phe
 100 105 110

Tyr Glu Leu Leu Thr Gly Gly Ser Tyr Glu Ser Gln Phe Ile Pro His
 115 120 125

Leu Phe Val His Asn Trp Ser Leu Ala Val Glu Val His Tyr Tyr Ile
 130 135 140

Leu Trp Gly Leu Ala Val Trp Phe Leu Ser Lys Gln Ala Lys Ser Asn
 145 150 155 160

Gly Gln Leu Lys Gly Met Val Phe Leu Leu Ser Ala Val Ala Phe Leu
 165 170 175

Ile Ser Phe Phe Ser Met Phe Ile Gly Ser Phe Leu Val Thr Ser Tyr
 180 185 190

Ser Ser Val Tyr Phe Ser Ser Leu Thr His Val Tyr Pro Phe Phe Leu
 195 200 205

Gly Ser Met Leu Ala Thr Ile Val Gly Val Arg Gln Thr Thr Ser Leu
 210 215 220

Val Lys Gln Leu Asp Lys Ile Trp Asp Leu Arg Lys Thr Leu Val Val
 225 230 235 240

Phe Gly Gly Gly Phe Gly Phe Leu Val Leu Leu Thr Phe Phe Val Lys
 245 250 255

Phe Thr Tyr Leu Phe Ala Tyr Leu Ile Gly Phe Leu Leu Ala Ser Leu
 260 265 270

Ala Ala Leu Ala Met Ile Leu Ala Ala Arg Val Leu His Glu Lys Thr
 275 280 285

His His Ile Gln Glu Ser Lys Ile Ile Ser Phe Leu Ala Asp Thr Ser
 290 295 300

Tyr Ala Val Tyr Leu Phe His Trp Pro Phe Tyr Ile Ile Phe Ser Gln
 305 310 315 320

Leu Thr Ser Asn Leu Leu Ala Val Leu Leu Thr Leu Ile Cys Ser Tyr
 325 330 335

Gly Phe Ala Ser Leu Ser Phe Tyr Val Leu Glu Pro Trp Ile Ala Gly
 340 345 350

Lys Asn Thr Pro Ile Val Gln Thr Leu Arg Pro Leu Pro Tyr Ile His
 355 360 365

Ala Ile Leu Ala Ala Gly Thr Gly Ile Leu Thr Ile Ile Val Cys Thr
 370 375 380

Val Thr Leu Leu Ala Pro Gln Val Gly Ala Phe Glu Thr Asp Leu Thr
 385 390 395 400

Val Asn Gly Leu Lys Gln Ala Ala Thr Asn Ile Gly Gln Thr Lys Val
 405 410 415

Met Ala Glu Arg Ala Asp Ala Asn Ser Leu Gly Ile Ala Asp Gly Thr
 420 425 430

Met Leu Ile Gly Asp Ser Val Ala Leu Arg Ala Asn Thr Ala Leu Gln
 435 440 445

Thr Ala Leu Pro Gly Ala Gln Ile Asn Ala Gln Val Ser Val Thr Thr
 450 455 460

Lys Thr Ala Asn Glu Ile Met Leu Asn Asn Ser Gln Asn Lys Phe Leu
 465 470 475 480

Pro Lys Thr Val Val Ile Ala Thr Gly Val Asn Asn Pro Glu Asn Tyr
 485 490 495

Lys Asp Asp Trp Asp Ser Ile Val Lys Asn Leu Pro Lys Gly His His
 500 505 510

Met Ile Leu Val Thr Pro Tyr Glu Gly Asp Lys Thr Lys Glu Thr Tyr
 515 520 525

Ala Ile Val Glu Lys Ala Ala Tyr Met Arg Glu Leu Ala Glu Lys
 530 535 540

Thr Pro Tyr Ile Thr Ile Ala Asp Trp Asn Gln Val Ala Lys Glu His
 545 550 555 560

Pro Glu Ile Trp Ala Gly Thr Asp Gln Val His Phe Gly Ser Glu Ser
 565 570 575

Ser Thr Ile Glu Ala Gly Ala Lys Leu Tyr Ala Asp Thr Ile Ala Thr
 580 585 590

Ala Leu Gln Thr Ala Gln Asp Lys Pro Val Lys Ser Lys
 595 600 605

<210> 396

<211> 119

<212> PRT

<213> Streptococcus pneumoniae

<400> 396

Met Ser Ile Ile Leu Thr Thr Ile Val Ala Leu Glu His Phe Tyr Ile
1 5 10 15

Phe Tyr Leu Glu Ser Ile Ala Thr Gln Ser Asp Ala Thr Ser Arg Val
20 25 30

Phe Asn Met Glu Lys Glu Glu Leu Ala His Pro Ser Val Ser Ser Leu
35 40 45

Phe Lys Asn Gln Gly Ile Tyr Lys Ala Leu Leu Gly Val Phe Leu Leu
50 55 60

Tyr Val Ile Tyr Phe Ser Gln Asn Leu Glu Ile Val Thr Ile Phe Val
65 70 75 80

Leu Phe Val Ile Gly Ala Ala Thr Tyr Gly Ser Leu Thr Ala Asp Lys
85 90 95

Lys Ile Ile Leu Lys Gln Gly Gly Ser Ala Ile Leu Ala Leu Ile Ser
100 105 110

Ile Leu Leu Phe Lys Tyr Thr
115

<210> 397

<211> 603

<212> PRT

<213> Streptococcus pneumoniae

<400> 397

Met Ala Val Ala Asn Cys Ala Lys Tyr Asn Ile Arg Arg Ser Lys Met
1 5 10 15

Lys Thr Val Gln Phe Phe Trp His Tyr Phe Lys Val Tyr Lys Phe Ser
20 25 30

Phe Val Val Val Ile Leu Met Ile Val Leu Ala Thr Phe Ala Gln Ala
35 40 45

Leu Phe Pro Val Phe Ser Gly Gln Ala Val Thr Gln Leu Ala Asn Leu
50 55 60

Val Gln Ala Tyr Gln Asn Gly Asn Pro Glu Leu Val Trp Gln Ser Leu
65 70 75 80

Ser Gly Ile Met Val Asn Leu Gly Leu Leu Val Leu Val Leu Phe Ile
85 90 95

Ser Ser Val Ile Tyr Met Cys Leu Met Thr Arg Val Ile Ala Glu Ser
100 105 110

Thr Asn Glu Met Arg Lys Gly Leu Phe Gly Lys Leu Ala Gln Leu Thr
115 120 125

Val Ser Phe Phe Asp Arg Arg Gln Asp Gly Asp Ile Leu Ser His Phe
130 135 140

Thr Ser Asp Leu Asp Asn Ile Leu Gln Ala Phe Asn Glu Ser Leu Ile
145 150 155 160

Gln Val Met Ser Asn Ile Val Leu Tyr Ile Gly Leu Ile Leu Val Met
165 170 175

Phe Ser Arg Asn Val Thr Leu Ala Leu Ile Thr Ile Ala Ser Thr Pro
180 185 190

Leu Ala Phe Leu Met Leu Ile Phe Ile Val Lys Met Ala Arg Lys Tyr
195 200 205

Thr Asn Leu Gln Gln Lys Glu Val Gly Lys Leu Asn Ala Tyr Met Asp
210 215 220

Glu Ser Ile Ser Gly Gln Lys Ala Val Ile Val Gln Gly Ile Gln Glu
225 230 235 240

Asp Met Met Ala Gly Phe Leu Glu Gln Asn Glu Arg Val Arg Lys Ala
245 250 255

Thr Phe Lys Gly Arg Met Phe Ser Gly Ile Leu Phe Pro Val Met Asn
260 265 270

Gly Met Ser Leu Ile Asn Thr Ala Ile Val Ile Phe Ala Gly Ser Ala
275 280 285

Val Leu Leu Asn Asp Lys Ser Ile Glu Thr Ser Thr Ala Leu Gly Leu
 290 295 300
 Ile Val Met Phe Ala Gln Phe Ser Gln Gln Tyr Tyr Gln Pro Ile Ile
 305 310 315 320
 Gln Val Ala Ala Ser Trp Gly Ser Leu Gln Leu Ala Phe Thr Gly Ala
 325 330 335
 Glu Arg Ile Gln Glu Met Phe Asp Ala Glu Glu Glu Ile Arg Pro Glu
 340 345 350
 Lys Ala Pro Thr Phe Thr Lys Leu Gln Glu Ser Val Glu Ile Ser His
 355 360 365
 Ile Val Phe Ser Tyr Leu Pro Asp Lys Pro Ile Leu Lys Asp Val Ser
 370 375 380
 Ile Ser Ala Pro Lys Gly Gln Met Thr Ala Val Val Gly Pro Thr Gly
 385 390 395 400
 Ser Gly Lys Thr Thr Ile Met Asn Leu Ile Asn Arg Phe Tyr Asp Val
 405 410 415
 Asp Ala Gly Gly Ile Tyr Phe Asp Gly Lys Asp Ile Arg Gly Tyr Asp
 420 425 430
 Leu Asp Ser Leu Arg Ser Lys Val Gly Ile Val Leu Gln Asp Ser Val
 435 440 445
 Leu Phe Ser Gly Thr Ile Arg Asp Asn Ile Arg Phe Gly Val Pro Asp
 450 455 460
 Ala Ser Gln Glu Met Val Glu Val Ala Ala Lys Ala Thr His Ile His
 465 470 475 480
 Asp Tyr Ile Glu Ser Leu Pro Asp Lys Tyr Asp Thr Leu Ile Asp Asp
 485 490 495
 Asp Gln Ser Ile Phe Ser Thr Gly Gln Lys Gln Leu Ile Ser Ile Ala

[illegible]

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      85              90              95

Ala Met Ser Ile Tyr Asp Asn Val Ala Tyr Gly Pro Arg Thr His Gly
      100              105              110

Ile Arg Asp Lys Lys Gln Leu Asp Ala Leu Val Glu Lys Ser Leu Lys
      115              120              125

Gly Ala Ala Ile Trp Glu Glu Val Lys Asp Asp Leu Lys Lys Ser Ala
      130              135              140

Met Ser Leu Ser Gly Gly Gln Gln Gln Arg Leu Cys Ile Ala Arg Ala
      145              150              155              160

Leu Ala Val Glu Pro Asp Ile Leu Leu Met Asp Glu Pro Thr Ser Ala
      165              170              175

Leu Asp Pro Ile Ser Thr Leu Lys Ile Glu Asp Leu Ile Gln Gln Leu
      180              185              190

Lys Lys Asp Tyr Thr Ile Ile Ile Val Thr His Asn Met Gln Gln Ala
      195              200              205

Ser Arg Ile Ser Asp Lys Thr Ala Phe Phe Leu Thr Gly Glu Ile Cys
      210              215              220

Glu Phe Gly Asp Thr Val Asp Val Phe Thr Asn Pro Lys Asp Gln Arg
      225              230              235              240

Thr Glu Asp Tyr Ile Ser Gly Arg Phe Gly
      245              250

<210> 399
<211> 226
<212> PRT
<213> Streptococcus pneumoniae

<400> 399

Met Met Lys Glu Ile Phe Asp Arg Arg Tyr Pro Val Thr Ser Phe Phe
1              5              10              15

Leu Leu Val Thr Ala Leu Val Phe Leu Leu Met Leu Val Thr Ala Gly

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	20		25		30	
Gly Asn Phe Asp Arg Ala Asp Thr Leu Phe Arg Phe Gly Ala Met Tyr	35		40		45	
Gly Pro Ala Ile Arg Leu Phe Pro Glu Gln Val Trp Arg Leu Leu Ser	50		55		60	
Ala Ile Phe Val His Ile Gly Trp Glu His Phe Ile Val Asn Met Leu	65		70		75	80
Ser Leu Tyr Tyr Leu Gly Arg Gln Val Glu Glu Ile Phe Gly Ser Lys		85		90		95
Gln Phe Phe Phe Leu Tyr Leu Leu Ser Gly Met Met Gly Asn Leu Phe		100		105		110
Val Phe Val Phe Ser Pro Lys Ser Leu Ala Ala Gly Ala Ser Thr Ser		115		120		125
Leu Tyr Gly Leu Phe Ala Ala Ile Ile Val Leu Arg Tyr Ala Thr Arg		130		135		140
Asn Pro Tyr Ile Gln Gln Leu Gly Gln Ser Tyr Leu Thr Leu Phe Val	145		150		155	160
Val Asn Ile Ile Gly Ser Val Leu Ile Pro Gly Ile Ser Leu Ala Gly		165		170		175
His Ile Gly Gly Ala Val Gly Gly Ala Phe Leu Ala Val Ile Phe Pro		180		185		190
Val Arg Gly Glu Lys Arg Met Tyr Asn Thr Ser Gln Arg Leu Gly Ala		195		200		205
Val Val Leu Phe Val Gly Leu Ala Ile Leu Leu Phe Tyr Lys Gly Met		210		215		220
Gly Leu	225					

<210> 400

<211> 232

<212> PRT

<213> Streptococcus pneumoniae

<400> 400

Met Thr Ala Thr Lys Met Asn Ala Gln Glu Ile Ile Gln Phe Ile Ala
 1 5 10 15

Asn Ala Glu Lys Lys Thr Ser Val Lys Val Thr Phe Glu Gly Gln Leu
 20 25 30

Ala Thr Ala Val Pro Ser Ser Val Val Lys Leu Gly Asn Val Leu Phe
 35 40 45

Gly Asp Trp Lys Asp Val Ala Pro Leu Leu Glu Gly Leu Val Glu Asn
 50 55 60

Gln Asp Tyr Val Val Glu Gln Asp Ala Arg Asn Ser Ala Val Pro Leu
 65 70 75 80

Leu Asp Lys Arg Ala Ile Asn Ala Arg Ile Glu Pro Gly Ala Ile Ile
 85 90 95

Arg Asp Gln Val Glu Ile Gly Asp Asn Ala Val Ile Met Met Gly Ser
 100 105 110

Val Ile Asn Ile Gly Ala Glu Ile Gly Ala Gly Thr Met Ile Asp Met
 115 120 125

Gly Ala Ile Leu Gly Gly Arg Ala Ile Val Gly Lys Asn Ser His Val
 130 135 140

Gly Ala Gly Ala Val Leu Ala Gly Val Ile Glu Pro Ala Ser Ala Glu
 145 150 155 160

Pro Val Arg Val Gly Asp Asn Val Leu Ile Gly Ala Asn Ala Val Val
 165 170 175

Ile Glu Gly Val Gln Ile Gly Ser Gly Ser Val Val Ala Ala Gly Ala
 180 185 190

Ile Val Thr Gln Asp Val Pro Glu Asn Val Val Val Ala Gly Val Pro
 195 200 205

Ala Arg Ile Ile Lys Glu Ile Asp Ala Gln Thr Gln Gln Lys Thr Ala
 210 215 220

Leu Glu Asp Ala Leu Arg Thr Leu
 225 230

<210> 401
 <211> 361
 <212> PRT
 <213> Streptococcus pneumoniae

<400> 401

Met Trp Gly Val Tyr Gln Leu His Pro Ala Pro Ala Val Gln Val Val
 1 5 10 15

Val Leu Ala Ile Ala Val Thr Gln Val Leu His Asp Leu Leu Leu Gln
 20 25 30

Gly Arg Asp Asp Asn Ser Leu Asn Glu Val Ala Asn Gln Trp Ile Ala
 35 40 45

Thr Phe Phe Leu Ser Cys Tyr Asn Lys Gly Met Asn Gln Tyr Gln Lys
 50 55 60

Lys Ile Val Asn Gly Lys Ile Tyr Ser Leu Leu Ser Gly Leu Ile Trp
 65 70 75 80

Gly Ile Cys Gly Ile Leu Gly Glu Tyr Phe Phe Thr His Tyr Gln Val
 85 90 95

Ser Ser Gly Trp Ile Thr Ser Met Arg Leu Thr Leu Ala Gly Ser Leu
 100 105 110

Val Leu Ile Trp Ser Ala Ile Gln Leu Lys Ser Gln Val Leu Asp Ile
 115 120 125

Trp Arg Asp Lys Lys Asn Tyr Leu Pro Phe Leu Ala Tyr Ala Ile Leu
 130 135 140

Gly Ile Phe Ser Val Gln Tyr Phe Phe Tyr Leu Cys Val Glu Tyr Ser
 145 150 155 160
 Asn Ala Thr Thr Ala Thr Ile Leu Gln Phe Ile Ser Pro Val Phe Ile
 165 170 175
 Leu Phe Tyr Asn Arg Leu Val Tyr Gln Lys Arg Ala Ser Lys Ser Ala
 180 185 190
 Val Phe Tyr Val Leu Val Ala Met Leu Gly Val Cys Leu Met Ala Thr
 195 200 205
 Lys Gly Asp Leu Ser Gln Leu Ser Met Thr Pro Leu Ala Leu Ile Thr
 210 215 220
 Gly Leu Leu Ser Ala Met Gly Val Met Phe Asn Val Ile Leu Pro Gln
 225 230 235 240
 Pro Phe Ala Lys Arg Tyr Gly Phe Val Pro Thr Val Gly Trp Gly Met
 245 250 255
 Ile Leu Ala Gly Leu Phe Ser Asn Val Leu Ser Pro Val Tyr Gln Leu
 260 265 270
 Ser Phe Thr Leu Asp Ile Trp Ser Ile Leu Ile Cys Leu Ile Ile Ala
 275 280 285
 Phe Phe Gly Thr Ala Phe Ala Phe Phe Ile Ser Met Lys Ala Val Ser
 290 295 300
 Leu Val Ser Pro Leu Val Val Ala Val Ile Ser Ala Ser Glu Pro Leu
 305 310 315 320
 Ser Ser Ala Leu Leu Ser Val Leu Phe Leu Gly Leu Val Val Asp Trp
 325 330 335
 Ser Leu Leu Leu Ala Ile Ala Leu Ile Ile Leu Pro Met Ile Phe Leu
 340 345 350
 Ser Ile Glu Glu Ala Lys Glu Ser Arg
 355 360

<210> 402
 <211> 190
 <212> PRT
 <213> Streptococcus pneumoniae

<400> 402

Met Asn Lys Leu Met Lys Phe Ile Ser Val Phe Leu Thr Ser Ile Val
 1 5 10 15

Leu Ile Val Ser Ala Ile Pro Ser Val Ser Ala Val Tyr Ala Ser Glu
 20 25 30

Gln Val Ser Gln Ile Glu Thr Asn Met Glu Leu Gln Pro Val Thr Ser
 35 40 45

Leu Thr Glu Glu Gln Ile Asn Thr Leu Ala Asn Glu Ile Gln Ser Phe
 50 55 60

His Pro Asp Val Ser Gln Gln Trp Ile Lys Glu Val Ile Asn Arg Gln
 65 70 75 80

Leu Gln Gly Asp Tyr Thr Ile Pro Pro Thr Tyr Ser Pro Phe Arg Ala
 85 90 95

Val Trp Gln Gly Ile Thr Val Asn Gln Met Gly Ala Leu Leu Asp Thr
 100 105 110

Ala Ile Ala Leu Ala Leu Gly Gly Thr Thr Ala Gly Leu Ala Asn Leu
 115 120 125

Ile Lys Val Lys Gly Lys His Ala Ala Lys Ser Ala Ile Arg Ser Ala
 130 135 140

Ile Ser Arg Tyr Leu Gly Ser Trp Phe Val Asn Asp Val Ala Leu Glu
 145 150 155 160

Phe Ala Met Asn Leu Leu Ser Pro Gly Thr Tyr Leu Ala Gln Leu Trp
 165 170 175

Asp Lys Asn Asp Ala Ile Pro Asn Asn Gly Arg Ile Asn Phe
 180 185 190

<210> 403
 <211> 435
 <212> PRT
 <213> Streptococcus pneumoniae

<400> 403

Met Lys Gly Val Asn Met Glu Lys Gln Gln Pro Ser Lys Ala Ala Leu
 1 5 10 15

Leu Ser Ile Ile Pro Gly Leu Gly Gln Ile Tyr Asn Lys Gln Lys Ala
 20 25 30

Lys Gly Phe Ile Phe Leu Gly Val Thr Ile Val Phe Val Leu Tyr Phe
 35 40 45

Leu Ala Leu Ala Thr Pro Glu Leu Ser Asn Leu Ile Thr Leu Gly Asp
 50 55 60

Lys Pro Gly Arg Asp Asn Ser Leu Phe Met Leu Ile Arg Gly Ala Phe
 65 70 75 80

His Leu Ile Phe Val Ile Val Tyr Val Leu Phe Tyr Phe Ser Asn Ile
 85 90 95

Lys Asp Ala His Thr Ile Ala Lys Arg Ile Asn Asn Gly Ile Pro Val
 100 105 110

Pro Arg Thr Leu Lys Asp Met Ile Lys Gly Ile Tyr Glu Asn Gly Phe
 115 120 125

Pro Tyr Leu Leu Ile Ile Pro Ser Tyr Val Ala Met Thr Phe Ala Ile
 130 135 140

Ile Phe Pro Val Ile Val Thr Leu Met Ile Ala Phe Thr Asn Tyr Asp
 145 150 155 160

Phe Gln His Leu Pro Pro Asn Lys Leu Leu Asp Trp Val Gly Leu Thr
 165 170 175

Asn Phe Thr Asn Ile Trp Ser Leu Ser Thr Phe Arg Ser Ala Phe Gly
 180 185 190

Ser Val Leu Ser Trp Thr Ile Ile Trp Ala Leu Ala Ala Ser Thr Leu
 195 200 205

Gln Ile Val Ile Gly Ile Phe Thr Ala Ile Ile Ala Asn Gln Pro Phe
 210 215 220

Ile Lys Gly Lys Arg Ile Phe Gly Val Ile Phe Leu Leu Pro Trp Ala
 225 230 235 240

Val Pro Ala Phe Ile Thr Ile Leu Thr Phe Ser Asn Met Phe Asn Asp
 245 250 255

Ser Val Gly Ala Ile Asn Thr Gln Val Leu Pro Ile Leu Ala Lys Phe
 260 265 270

Leu Pro Phe Leu Asp Gly Ala Leu Ile Pro Trp Lys Thr Asp Pro Thr
 275 280 285

Trp Thr Lys Ile Ala Leu Ile Met Met Gln Gly Trp Leu Gly Phe Pro
 290 295 300

Tyr Ile Tyr Val Leu Thr Leu Gly Ile Leu Gln Ser Ile Pro Asn Asp
 305 310 315 320

Leu Tyr Glu Ala Ala Tyr Ile Asp Gly Ala Asn Ala Trp Gln Lys Phe
 325 330 335

Arg Asn Ile Thr Phe Pro Met Ile Leu Ala Val Ala Ala Pro Thr Leu
 340 345 350

Ile Ser Gln Tyr Thr Phe Asn Phe Asn Asn Phe Ser Ile Met Tyr Leu
 355 360 365

Phe Asn Gly Gly Gly Pro Gly Ser Val Gly Gly Gly Ala Gly Ser Thr
 370 375 380

Asp Ile Leu Ile Ser Trp Ile Tyr Arg Leu Thr Thr Gly Thr Ser Pro
 385 390 395 400

Gln Tyr Ser Met Ala Ala Ala Val Thr Leu Ile Ile Ser Ile Ile Val
 405 410 415

Ile Ser Ile Ser Met Ile Ala Phe Lys Lys Leu His Ala Phe Asp Met
 420 425 430

Glu Asp Val
 435

<210> 404
 <211> 287
 <212> PRT
 <213> Streptococcus pneumoniae
 <400> 404

Met Ile Trp Arg Thr Ser Lys Met Asn Asn Ser Ile Lys Leu Lys Arg
 1 5 10 15

Arg Leu Thr Gln Ser Leu Thr Tyr Leu Tyr Leu Ile Gly Leu Ser Ile
 20 25 30

Val Ile Ile Tyr Pro Leu Leu Ile Thr Ile Met Ser Ala Phe Lys Ala
 35 40 45

Gly Asn Val Ser Ala Phe Lys Leu Asp Thr Asn Ile Asp Leu Asn Phe
 50 55 60

Asp Asn Phe Lys Gly Leu Phe Thr Glu Thr Leu Tyr Gly Thr Trp Tyr
 65 70 75 80

Leu Asn Thr Leu Ile Ile Ala Leu Ile Thr Met Ala Val Gln Thr Ser
 85 90 95

Ile Ile Val Leu Ala Gly Tyr Ala Tyr Ser Arg Tyr Asn Phe Leu Ala
 100 105 110

Arg Lys Gln Ser Leu Val Phe Phe Leu Ile Ile Gln Met Val Pro Thr
 115 120 125

Met Ala Ala Leu Thr Ala Phe Phe Val Met Ala Leu Met Leu Asn Ala
 130 135 140

Leu Asn His Asn Trp Phe Leu Ile Phe Leu Tyr Val Gly Gly Gly Ile
 145 150 155 160

Pro Met Asn Ala Trp Leu Met Lys Gly Tyr Phe Asp Thr Val Pro Met
165 170 175

Ser Leu Asp Glu Ser Ala Lys Leu Asp Gly Ala Gly His Phe Arg Arg
180 185 190

Phe Trp Gln Ile Val Leu Pro Leu Val Arg Pro Met Val Ala Val Gln
195 200 205

Ala Leu Trp Ala Phe Met Gly Pro Phe Gly Asp Tyr Ile Leu Ser Ser
210 215 220

Phe Leu Leu Arg Glu Lys Glu Tyr Phe Thr Val Ala Val Gly Leu Gln
225 230 235 240

Thr Phe Val Asn Asn Ala Lys Asn Leu Lys Ile Ala Tyr Phe Ser Ala
245 250 255

Gly Ala Ile Leu Ile Ala Leu Pro Ile Cys Ile Leu Phe Phe Phe Leu
260 265 270

Gln Lys Asn Phe Val Ser Gly Leu Thr Ser Gly Gly Asp Lys Gly
275 280 285

<210> 405

<211> 266

<212> PRT

<213> Streptococcus pneumoniae

<400> 405

Met Leu Pro Tyr Pro Phe Ser Tyr Phe Ser Ser Ile Trp Gly Val Arg
1 5 10 15

Lys Pro Leu Ser Lys Arg Phe Glu Leu Asn Trp Phe Gln Leu Leu Phe
20 25 30

Thr Ser Ile Phe Leu Ile Ser Leu Ser Met Val Pro Ile Ala Ile Gln
35 40 45

Asn Ser Ser Gln Glu Thr Tyr Pro Leu Glu Thr Phe Ile Asp Asn Val
50 55 60

Tyr Glu Pro Leu Thr Asp Lys Val Val Gln Asp Leu Ser Glu His Ala
 65 70 75 80
 Thr Ile Val Asp Gly Thr Leu Thr Tyr Thr Gly Thr Ala Ser Gln Ala
 85 90 95
 Pro Ser Val Val Ile Gly Pro Ser Gln Ile Lys Glu Leu Pro Lys Asp
 100 105 110
 Leu Gln Leu His Phe Asp Thr Asn Glu Leu Val Ile Ser Lys Glu Ser
 115 120 125
 Lys Glu Leu Thr Arg Ile Ser Tyr Arg Ala Ile Gln Thr Glu Ser Phe
 130 135 140
 Lys Ser Lys Asp Ser Leu Thr Gln Ala Ile Ser Lys Asp Trp Tyr Gln
 145 150 155 160
 Gln Asn Arg Val Tyr Ile Ser Leu Phe Leu Val Leu Gly Ala Ser Phe
 165 170 175
 Leu Phe Gly Leu Asn Phe Phe Ile Val Ser Leu Gly Ala Ser Phe Leu
 180 185 190
 Leu Tyr Ile Thr Lys Arg Ser Arg Leu Phe Ser Phe Asn Thr Phe Lys
 195 200 205
 Glu Cys Tyr His Phe Ile Leu Asn Cys Leu Gly Leu Pro Thr Leu Ile
 210 215 220
 Thr Leu Ile Leu Gly Leu Phe Gly Gln Asn Met Thr Thr Leu Ile Thr
 225 230 235 240
 Val Gln Asn Ile Leu Phe Val Leu Tyr Leu Val Thr Ile Phe Tyr Lys
 245 250 255
 Thr His Phe Arg Asp Pro Asn Tyr His Lys
 260 265
 <210> 406

<211> 313
 <212> PRT
 <213> Streptococcus pneumoniae
 <400> 406

Met Lys Gln Thr Lys Arg Ile Lys Arg Trp Arg Tyr Tyr Leu Arg Arg
 1 5 10 15

Phe Ala Tyr Gln Ile Lys Ile Leu Arg Val Leu Gln Ser Ile Ser Arg
 20 25 30

Glu Lys Tyr Asp Glu Lys Ile Ser Ala Ser Leu Val Tyr Gly Phe Leu
 35 40 45

Ser Ala Val Ala Val Asn Phe Phe Gln Pro Gly His Val Tyr Ser
 50 55 60

Ser Gly Ala Thr Gly Leu Ala Gln Ile Ile Ser Ala Leu Ser Asn His
 65 70 75 80

Trp Phe Gly Phe His Ile Pro Ile Ser Leu Ser Phe Tyr Ala Ile Asn
 85 90 95

Phe Pro Leu Met Val Leu Ala Trp Tyr Gln Ile Gly His Lys Phe Thr
 100 105 110

Val Phe Thr Phe Ile Thr Val Ser Met Ser Ser Phe Phe Ile Gln Phe
 115 120 125

Val Pro Val Ala Thr Leu Thr Glu Asp Pro Ile Ile Asn Ser Leu Phe
 130 135 140

Gly Gly Val Val Met Gly Leu Gly Ile Gly Phe Ala Leu Arg Asn Asn
 145 150 155 160

Ile Ser Ser Gly Gly Thr Asp Ile Val Ser Leu Thr Ile Arg Lys Lys
 165 170 175

Thr Gly Lys Asn Val Gly Ser Ile Ser Phe Leu Val Asn Gly Thr Ile
 180 185 190

Met Leu Ile Ala Gly Leu Thr Phe Gly Trp Lys Tyr Ala Leu Tyr Ser

195 200 205
 Met Ile Thr Ile Phe Val Ser Ser Arg Val Thr Asp Ala Val Phe Thr
 210 215 220
 Lys Gln Lys Arg Met Gln Ala Met Ile Val Thr Asn His Pro Glu Lys
 225 230 235 240
 Val Ile Glu Lys Ile His Lys Lys Leu His Arg Gly Ala Thr Met Ile
 245 250 255
 His Asp Ala Glu Gly Thr Tyr Asn His Glu Arg Lys Ala Val Leu Ile
 260 265 270
 Thr Val Ile Thr Arg Ala Glu Phe Asn Glu Phe Lys Gln Ile Met Thr
 275 280 285
 Gln Val Asp Pro Ser Ser Phe Val Ser Val Ser Glu Asn Val His Ile
 290 295 300
 Leu Gly Arg Phe Val Glu Ile Asp Asn
 305 310
 <210> 407
 <211> 76
 <212> PRT
 <213> Streptococcus pneumoniae
 <400> 407
 Met Lys Arg Val Ile Leu Leu Ala Val Ile Gln Ala Val Val Leu Phe
 1 5 10 15
 Phe Ile Ile Gly Ala Leu Ala Tyr Ala Phe Lys Gly Asp Phe Phe Tyr
 20 25 30
 Asn Tyr Leu Ala Val Val Phe Ala Pro Ile Ala Gly Val Leu Arg Phe
 35 40 45
 Gly Thr Ala Tyr Ile Thr Glu Ile Val Leu Pro Arg Lys Ala Ala Glu
 50 55 60
 Ile Ala Glu Lys Arg Lys Ala Gly Lys Asn Ser Lys

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65
70
75

<210> 408
<211> 213
<212> PRT
<213> Streptococcus pneumoniae

<400> 408

Met Lys Leu Leu Lys Asn Leu Gly Trp Ile Leu Leu Ala Leu Leu Ser
1 5 10 15

Phe Leu Phe Ile Tyr Gly Phe Ile Gln Gly Leu Ala Thr Ala Ser Leu
20 25 30

Ala Leu Gly Ala Ser Pro Tyr Ala Val Thr Leu Leu Tyr Val Ala Leu
35 40 45

Ala Gly Val Tyr Val Tyr Gly Ile Tyr Lys Trp Tyr Gln Lys Ala Pro
50 55 60

Val His Ile Glu Lys Ser Gly Phe Asn Arg Phe Ile Trp Leu Pro Val
65 70 75 80

Leu Val Trp Phe Leu Ser Leu Val Val Gln Phe Phe Leu Pro Asp Asp
85 90 95

Pro Ser Val Asn Gln Gln Ile Ala Thr Asp Leu Thr Leu Ser Gln Pro
100 105 110

Leu Phe Ser Phe Phe Ala Val Val Ile Phe Ala Pro Leu Thr Glu Glu
115 120 125

Ile Val Phe Arg Gly Met Leu Ala Arg Tyr Leu Phe Pro Lys Gln Asp
130 135 140

Asn Ser Lys Arg Thr Leu Ile Phe Leu Leu Val Ser Ser Leu Leu Phe
145 150 155 160

Ala Leu Ile His Phe Pro Gly Asp Val Gln Gln Phe Phe Val Tyr Phe
165 170 175

Ser Leu Gly Phe Ser Leu Gly Leu Ala Tyr Ile Ser Arg Lys Gly Leu

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180 185 190
 Val Tyr Ser Ile Ser Leu His Ala Leu Asn Asn Leu Val Gly Phe Leu
 195 200 205

 Met Ile Leu Met Leu
 210

 <210> 409
 <211> 226
 <212> PRT
 <213> Streptococcus pneumoniae

 <400> 409

 Met Gly Met Phe Val Gly Met Phe Lys Ala Arg Val Glu Ser His Glu
 1 5 10 15

 Ile Ile Leu Asp Val Lys Ala Leu Met Pro Trp Ile Ser Ala Ile Cys
 20 25 30

 Leu Leu Ile Gly Phe Ile Ser Met Phe Leu Thr Phe Asn Phe Leu Lys
 35 40 45

 Lys Ser Arg Lys Phe His Ser Leu Tyr Gln Glu Glu Met Asp Asp Asp
 50 55 60

 Leu Asn Glu Thr Tyr Tyr Val Gln Met Tyr Arg Asn Leu Glu Phe Gly
 65 70 75 80

 Thr Ile Ala Phe Asn Ile Thr Gly Val Ala Ile Pro Leu Ala Ile Phe
 85 90 95

 Ile Ser Leu Ser Glu Val Ile Ile Leu His Thr Asn Pro Gln Thr Phe
 100 105 110

 Phe Leu Ser Phe Leu Leu Phe Val Val Phe Leu Val Ala Gln Lys Ser
 115 120 125

 Leu Phe Lys Thr Ile Ala Ile Val Arg Gln Phe Asp Leu Glu Phe Phe
 130 135 140

 Ala Thr Pro Lys Asp Val Leu Asn Tyr Ile Asn Ser Tyr Asp Glu Gly

145 150 155 160
 Glu Arg Gln Ala Asn Leu Glu Gln Ser Phe Arg Ile Leu Phe Gln Leu
 165 170 175
 His Gln Tyr Val Leu Pro Ala Leu Tyr Ile Phe Leu Ile Ile Ile Ser
 180 185 190
 Phe Leu Thr Gly Glu Ile Gln Leu Leu Ala Phe Leu Leu Val Gly Ala
 195 200 205
 Ile His Val Tyr Ile Asn Val Met Gln Leu Pro Met Val Lys Arg Tyr
 210 215 220
 Phe Lys
 225
 <210> 410
 <211> 621
 <212> PRT
 <213> Streptococcus pneumoniae
 <400> 410
 Met Lys Lys Thr Thr Ile Leu Ser Leu Thr Thr Ala Ala Val Ile Leu
 1 5 10 15
 Ala Ala Tyr Val Pro Asn Glu Pro Ile Leu Ala Asp Thr Pro Ser Ser
 20 25 30
 Glu Val Ile Lys Glu Thr Lys Val Gly Ser Ile Ile Gln Gln Asn Asn
 35 40 45
 Ile Lys Tyr Lys Val Leu Thr Val Glu Gly Asn Ile Gly Thr Val Gln
 50 55 60
 Val Gly Asn Gly Val Thr Pro Val Glu Phe Glu Ala Gly Gln Asp Gly
 65 70 75 80
 Lys Pro Phe Thr Ile Pro Thr Lys Ile Thr Val Gly Asp Lys Val Phe
 85 90 95
 Thr Val Thr Glu Val Ala Ser Gln Ala Phe Ser Tyr Tyr Pro Asp Glu

100	105	110
Thr Gly Arg Ile Val Tyr Tyr Pro Ser Ser Ile Thr Ile Pro Ser Ser 115 120		
Ile Lys Lys Ile Gln Lys Lys Gly Phe His Gly Ser Lys Ala Lys Thr 130 135 140		
Ile Ile Phe Asp Lys Gly Ser Gln Leu Glu Lys Ile Glu Asp Arg Ala 145 150 155 160		
Phe Asp Phe Ser Glu Leu Glu Glu Ile Glu Leu Pro Ala Ser Leu Glu 165 170 175		
Tyr Ile Gly Thr Ser Ala Phe Ser Phe Ser Gln Lys Leu Lys Lys Leu 180 185 190		
Thr Phe Ser Ser Ser Ser Lys Leu Glu Leu Ile Ser His Glu Ala Phe 195 200 205		
Ala Asn Leu Ser Asn Leu Glu Lys Leu Thr Leu Pro Lys Ser Val Lys 210 215 220		
Thr Leu Gly Ser Asn Leu Phe Arg Leu Thr Thr Ser Leu Lys His Val 225 230 235 240		
Asp Val Glu Glu Gly Asn Glu Ser Phe Ala Ser Val Asp Gly Val Leu 245 250 255		
Phe Ser Lys Asp Lys Thr Gln Leu Ile Tyr Tyr Pro Ser Gln Lys Asn 260 265 270		
Asp Glu Ser Tyr Lys Thr Pro Lys Glu Thr Lys Glu Leu Ala Ser Tyr 275 280 285		
Ser Phe Asn Lys Asn Ser Tyr Leu Lys Lys Leu Glu Leu Asn Glu Gly 290 295 300		
Leu Glu Lys Ile Gly Thr Phe Ala Phe Ala Asp Ala Ile Lys Leu Glu 305 310 315 320		

Glu Ile Ser Leu Pro Asn Ser Leu Glu Thr Ile Glu Arg Leu Ala Phe
 325 330 335

Tyr Gly Asn Leu Glu Leu Lys Glu Leu Ile Leu Pro Asp Asn Val Lys
 340 345 350

Asn Phe Gly Lys His Val Met Asn Gly Leu Pro Lys Leu Lys Ser Leu
 355 360 365

Thr Ile Gly Asn Asn Ile Asn Ser Leu Pro Ser Phe Phe Leu Ser Gly
 370 375 380

Val Leu Asp Ser Leu Lys Glu Ile His Ile Lys Asn Lys Ser Thr Glu
 385 390 395 400

Phe Ser Val Lys Lys Asp Thr Phe Ala Ile Pro Glu Thr Val Lys Phe
 405 410 415

Tyr Val Thr Ser Glu His Ile Lys Asp Val Leu Lys Ser Asn Leu Ser
 420 425 430

Thr Ser Asn Asp Ile Ile Val Glu Lys Val Asp Asn Ile Lys Gln Glu
 435 440 445

Thr Asp Val Ala Lys Pro Lys Lys Asn Ser Asn Gln Gly Val Val Gly
 450 455 460

Trp Val Lys Asp Lys Gly Leu Trp Tyr Tyr Leu Asn Glu Ser Gly Ser
 465 470 475 480

Met Ala Thr Gly Trp Val Lys Asp Lys Gly Leu Trp Tyr Tyr Leu Asn
 485 490 495

Glu Ser Gly Ser Met Ala Thr Gly Trp Val Lys Asp Lys Gly Leu Trp
 500 505 510

Tyr Tyr Leu Asn Glu Ser Gly Ser Met Ala Thr Gly Trp Val Lys Asp
 515 520 525

Lys Gly Leu Trp Tyr Tyr Leu Asn Glu Ser Gly Ser Met Ala Thr Gly
 530 535 540

Trp Val Lys Asp Lys Gly Leu Trp Tyr Tyr Leu Asn Glu Ser Gly Ser
545 550 555 560

Met Ala Thr Gly Trp Val Lys Asp Lys Gly Leu Trp Tyr Tyr Leu Asn
565 570 575

Glu Ser Gly Ser Met Ala Thr Gly Trp Val Thr Val Ser Gly Lys Trp
580 585 590

Tyr Tyr Thr Tyr Asn Ser Gly Asp Leu Leu Val Asn Thr Thr Thr Pro
595 600 605

Asp Gly Tyr Arg Val Asn Ala Asn Gly Glu Trp Val Gly
610 615 620

<210> 411

<211> 383

<212> PRT

<213> Streptococcus pneumoniae

<400> 411

Met Ala Ile Phe Tyr Val Pro Ala Val Asn Leu Ile Gly Lys Gly Val
1 5 10 15

Val Asn Glu Val Gly Pro Tyr Ile Lys Glu Leu Gly Tyr Lys Lys Ala
20 25 30

Leu Leu Val Thr Asp Lys Tyr Ile Glu Gly Ser Asp Ile Leu Pro Lys
35 40 45

Thr Leu Lys Pro Leu Asp Thr Glu Gly Ile Glu Tyr Val Ile Phe Ser
50 55 60

Asp Val Glu Pro Asn Pro Thr Cys Lys Asn Val Thr Asp Gly Val Ala
65 70 75 80

Ala Leu Gln Glu His Gly Cys Asp Phe Ile Ile Ser Leu Gly Gly Gly
85 90 95

Ser Pro Gln Asp Ala Ala Ser Cys Ile Ser Ile Met Ala Thr Asn Gly
100 105 110

Gly Lys Pro Gln Asp Tyr Glu Gly Leu His Lys Ser Ala Lys Lys Gly
 115 120 125

Leu Pro Val Val Ala Ile Asn Thr Thr Ala Gly Thr Ser Ala Glu Ile
 130 135 140

Thr Ile Asn Tyr Val Ile Thr Asp Glu Glu Arg Lys Val Lys Met Val
 145 150 155 160

Met Val Asp Lys Asn Ser Leu Ala Leu Ile Ser Val Asn Asp Pro Glu
 165 170 175

Leu Met Leu Ser Lys Pro Lys Gly Leu Thr Ala Ala Thr Gly Met Asp
 180 185 190

Ala Leu Thr His Ala Val Glu Ala Leu Val Thr Pro Gly Ala Tyr Asp
 195 200 205

Val Thr Lys Lys Leu Ser Ile Gly Ala Ile Glu Leu Ile Lys Glu Tyr
 210 215 220

Leu Pro Arg Ala Val Glu Asn Gly His Asp Ile Glu Ala Arg Glu Gly
 225 230 235 240

Met Val Asn Ala Ile Phe Leu Gly Gly Met Ser Phe Asn Asn Ala Gly
 245 250 255

Leu Gly Tyr Val His Ser Met Ala His Gln Leu Gly Ala Val Tyr Asn
 260 265 270

Leu Pro His Gly Val Cys Cys Ala Met Leu Leu Pro Val Ile Glu Arg
 275 280 285

Glu Asn Ala Lys Arg Val Pro Glu Ala Phe Arg Asn Val Ala Lys Ala
 290 295 300

Leu Gly Leu His Val Glu Gly Lys Ser Asp Gln Glu Cys Ala Asp Tyr
 305 310 315 320

Ala Ile Ala Glu Ile Glu Lys Leu Ser Glu Thr Val Gly Ile Pro Lys
 325 330 335

Lys Leu Thr Glu Leu Gly Ile Glu Glu Lys Asp Phe Asp Phe Glu Tyr
 340 345 350

Leu Ser Lys Asn Ala Leu Ile Asp Ala Cys Ala Pro Gly Asn Pro Phe
 355 360 365

Met Pro Thr Leu Glu Glu Thr Ile Ala Phe Tyr Lys Glu Leu Phe
 370 375 380

<210> 412

<211> 257

<212> PRT

<213> Streptococcus pneumoniae

<400> 412

Met Ile Lys Asp Glu Arg Val Leu Glu Leu Ile Glu Ile Ile Lys Lys
 1 5 10 15

Lys Lys Arg Ile Ala Val Lys Glu Leu Ala Glu Ile Thr Phe Ser Ser
 20 25 30

Thr Ser Thr Leu Arg Arg Asp Leu Ile Phe Leu Glu Asn Gln Gly Leu
 35 40 45

Ile Lys Arg Lys His Gly Tyr Val Thr Leu Ser Ser Met Asn Thr Ile
 50 55 60

Glu Leu Ser His Gln Ile Arg Glu Gly Glu Ser Thr Arg Gln Lys Arg
 65 70 75 80

Leu Ile Ala Ser Leu Ala Lys Asp Phe Ile Arg Ser Gly Met Cys Ile
 85 90 95

Tyr Leu Asp Ser Ser Thr Thr Val Tyr Glu Leu Cys Pro Tyr Leu Ser
 100 105 110

Glu Leu Asp Asn Leu Ile Ile Phe Thr Asn Gly Leu His Thr Ala Gln
 115 120 125

Thr Leu Ser Glu Thr Val Lys Asp Ser Ser Lys Ile Phe Ile Thr Ser
 130 135 140

Gly Glu Val Lys His Gln Ser Cys Ser Val Val Asn Tyr Asp Lys Glu
 145 150 155 160

Asn Ser Leu Leu Asp His Phe Asn Ile Asp Leu Ala Phe Cys Ser Ala
 165 170 175

Arg Gly Ile Asp Asp Gln Tyr Val Tyr Glu Ala Ser Leu Ser Gln Ala
 180 185 190

Ile Ser Lys Lys Asn Ile Ile Asp Lys Ala His Glu Thr Ile Leu Leu
 195 200 205

Ile Asp Ser Ser Lys Phe Tyr Lys Thr Gly Phe Phe Lys Ile Asn Pro
 210 215 220

Leu Ser Lys Tyr Thr Thr Phe Ile Ser Asp Thr Val Pro Asp Gln Lys
 225 230 235 240

Leu Leu Asp Ala Val Glu Leu Phe Asp Gly Glu Trp Val Ser Asp Ile
 245 250 255

Gln

<210> 413
 <211> 268
 <212> PRT
 <213> Streptococcus pneumoniae

<400> 413

Met Leu Ser Leu Leu Ser Tyr Asp Phe Ile Gln Arg Ala Phe Leu Ala
 1 5 10 15

Val Ile Ala Met Ser Leu Phe Ser Pro Val Leu Gly Thr Phe Leu Ile
 20 25 30

Leu Arg Arg Gln Ser Leu Met Ser Asp Thr Leu Ser His Val Ser Leu
 35 40 45

Ser Gly Val Ala Phe Gly Leu Val Leu Gly Ile Ser Pro Thr Val Ser
 50 55 60

Thr Ile Ala Ile Val Leu Ile Ala Ala Val Phe Leu Glu Tyr Leu Arg
 65 70 75 80

Thr Val Tyr Lys Ser Phe Met Glu Ile Gly Thr Ala Ile Leu Met Ser
 85 90 95

Thr Gly Leu Ala Val Ser Leu Ile Val Met Ser Lys Gly Lys Ser Ser
 100 105 110

Ser Ser Met Ser Leu Asp Gln Tyr Leu Phe Gly Ser Ile Val Thr Ile
 115 120 125

Ser Glu Glu Gln Val Ile Ser Leu Phe Val Ile Ala Ala Val Val Leu
 130 135 140

Ile Leu Thr Phe Leu Phe Leu Arg Pro Met Tyr Ile Leu Thr Phe Asp
 145 150 155 160

Glu Asp Thr Ala Phe Val Asp Gly Leu Pro Val Arg Thr Met Ser Ile
 165 170 175

Leu Phe Asn Met Val Thr Gly Val Ala Ile Ala Leu Met Ile Pro Ala
 180 185 190

Ala Gly Ala Leu Leu Val Ser Thr Ile Met Val Leu Pro Ala Ser Ile
 195 200 205

Ala Leu Arg Leu Gly Lys Asn Phe Lys Ser Val Met Leu Leu Ala Ser
 210 215 220

Ala Ile Gly Phe Leu Gly Met Val Ala Gly Leu Tyr Ile Ser Tyr Tyr
 225 230 235 240

Ala Glu Thr Pro Ala Ser Ala Ser Ile Thr Ile Ile Phe Val Thr Val
 245 250 255

Phe Ile Leu Ile Ser Leu Val Arg Arg Phe Ile Lys
 260 265

<210> 414

<211> 234

<212> PRT

<213> Streptococcus pneumoniae

<400> 414

Met Arg Tyr Ile Thr Val Glu Asp Leu Ser Phe Tyr Tyr Asp Lys Glu
 1 5 10 15

Pro Val Leu Glu His Ile Asn Tyr Cys Val Asp Ser Gly Glu Phe Val
 20 25 30

Thr Leu Thr Gly Glu Asn Gly Ala Ala Lys Thr Thr Leu Ile Lys Ala
 35 40 45

Ser Leu Gly Ile Leu Gln Pro Arg Ile Gly Lys Val Ala Ile Ser Lys
 50 55 60

Thr Asn Thr Gln Gly Lys Lys Leu Arg Ile Ala Tyr Leu Pro Gln Gln
 65 70 75 80

Ile Ala Ser Phe Asn Ala Gly Phe Pro Ser Thr Val Tyr Glu Phe Val
 85 90 95

Lys Ser Gly Arg Tyr Pro Arg Lys Gly Trp Phe Arg Arg Leu Asn Ala
 100 105 110

His Asp Glu Glu His Ile Lys Ala Ser Leu Asp Ser Val Gly Met Trp
 115 120 125

Glu His Arg Asp Lys Arg Leu Gly Ser Leu Ser Gly Gly Gln Lys Gln
 130 135 140

Arg Ala Val Ile Ala Arg Met Phe Ala Ser Asp Pro Asp Val Phe Ile
 145 150 155 160

Leu Asp Glu Pro Thr Thr Gly Met Asp Ala Gly Ser Lys Asn Glu Phe
 165 170 175

Tyr Glu Leu Met His His Ser Ala His His His Gly Lys Ala Val Leu
 180 185 190

Met Ile Thr His Asp Pro Glu Glu Val Lys Asp Tyr Ala Asp Arg Asn

195 200 205
 Ile His Leu Val Arg Asn Gln Asp Ser Pro Trp Arg Cys Phe Asn Val
 210 215 220
 His Glu Asn Gly Gln Glu Val Gly His Ala
 225 230
 <210> 415
 <211> 516
 <212> PRT
 <213> Streptococcus pneumoniae
 <400> 415
 Met Ser Asn Lys Pro Ile Ala Asp Met Ile Glu Thr Ile Glu His Phe
 1 5 10 15
 Ala Gln Thr Gln Pro Ser Tyr Pro Val Tyr Asn Val Leu Gly Gln Glu
 20 25 30
 His Thr Tyr Gly Asp Leu Lys Ala Asp Ser Asp Ser Leu Ala Ala Val
 35 40 45
 Ile Asp Gln Leu Gly Leu Pro Glu Lys Ser Pro Val Val Val Phe Gly
 50 55 60
 Gly Gln Glu Tyr Glu Met Leu Ala Thr Phe Val Ala Leu Thr Lys Ser
 65 70 75 80
 Gly His Ala Tyr Ile Pro Ile Asp Ser His Ser Ala Leu Glu Arg Val
 85 90 95
 Ser Ala Ile Leu Glu Val Ala Glu Pro Ser Leu Ile Ile Ala Ile Ser
 100 105 110
 Ala Phe Pro Leu Glu Gln Val Ser Thr Pro Met Ile Asn Leu Ala Gln
 115 120 125
 Val Gln Glu Ala Phe Ala Gln Gly Asn Asn Tyr Glu Ile Thr His Pro
 130 135 140
 Val Lys Gly Asp Asp Asn Tyr Tyr Ile Ile Phe Thr Ser Gly Thr Thr

145		150		155		160
Gly Lys Pro Lys	Gly Val Gln Ile Ser His Asp Asn Leu Leu Ser Phe					
	165			170		175
Thr Asn Trp Met	Ile Thr Asp Lys Glu Phe Ala Thr Pro Ser Arg Pro					
	180		185			190
Gln Met Leu Ala Gln Pro Pro Tyr Ser Phe Asp Leu Ser Val Met Tyr						
	195		200			205
Trp Ala Pro Thr Leu Ala Leu Gly Gly Thr Leu Phe Thr Leu Pro Ser						
	210		215			220
Val Ile Thr Gln Asp Phe Lys Gln Leu Phe Ala Ala Ile Phe Ser Leu						
	225		230		235	240
Pro Ile Ala Ile Trp Thr Ser Thr Pro Ser Phe Ala Asp Met Ala Met						
	245			250		255
Leu Ser Glu Tyr Phe Asn Ser Glu Lys Met Pro Gly Ile Thr His Phe						
	260		265			270
Tyr Phe Asp Gly Glu Glu Leu Thr Val Lys Thr Ala Gln Lys Leu Arg						
	275		280			285
Glu Arg Phe Pro Asn Ala Arg Ile Ile Asn Ala Tyr Gly Pro Thr Glu						
	290		295			300
Ala Thr Val Ala Leu Ser Ala Val Ala Val Thr Asp Glu Met Leu Ala						
	305		310		315	320
Thr Leu Lys Arg Leu Pro Ile Gly Tyr Thr Lys Ala Asp Ser Pro Thr						
	325			330		335
Phe Ile Ile Asp Glu Glu Gly Asn Lys Leu Pro Asn Gly Glu Gln Gly						
	340		345			350
Glu Ile Ile Val Ser Gly Pro Ala Val Ser Lys Gly Tyr Met Asn Asn						
	355		360			365

Pro Glu Lys Thr Ala Glu Ala Phe Phe Glu Phe Glu Asp Leu Pro Ala
 370 375 380

Tyr His Thr Gly Asp Val Gly Thr Met Thr Asp Glu Gly Leu Leu Leu
 385 390 395 400

Tyr Gly Gly Arg Met Asp Phe Gln Ile Lys Phe Asn Gly Tyr Arg Ile
 405 410 415

Glu Leu Glu Asp Val Ser Gln Asn Leu Asn Lys Ser Arg Phe Ile Glu
 420 425 430

Ser Ala Val Ala Val Pro Arg Tyr Asn Lys Asp His Lys Val Gln Asn
 435 440 445

Leu Leu Ala Tyr Val Ile Leu Lys Asp Gly Val Arg Glu Gln Phe Glu
 450 455 460

Arg Asp Ile Asp Ile Thr Lys Ala Ile Lys Glu Asp Leu Thr Asp Ile
 465 470 475 480

Met Met Ser Tyr Met Met Pro Ser Lys Phe Leu Tyr Arg Asp Ser Leu
 485 490 495

Pro Leu Thr Pro Asn Gly Lys Ile Asp Ile Lys Gly Leu Ile Asn Glu
 500 505 510

Val Asn Lys Arg
 515

<210> 416

<211> 131

<212> PRT

<213> Streptococcus pneumoniae

<400> 416

Met Met Ser Leu Val His Asn Ile Ile Glu Asp Gly Asp Thr Glu Ser
 1 5 10 15

Leu Ile Arg Lys Tyr Leu His Ser Gly Val Ile Ile Asn Gly Gln Arg
 20 25 30

Tyr Lys Thr Leu Val Gly Thr Pro Gln Gly Gly Asn Leu Ser Pro Leu
 35 40 45

Leu Ser Asn Ile Met Leu Asn Glu Leu Asp Lys Glu Leu Glu Lys Arg
 50 55 60

Gly Leu Arg Phe Val Arg Tyr Ala Asp Asp Cys Val Ile Thr Val Gly
 65 70 75 80

Ser Glu Ala Ala Ser Lys Arg Val Met Tyr Ser Val Ser Arg Phe Ile
 85 90 95

Glu Lys Arg Leu Gly Leu Lys Val Asn Met Thr Lys Arg Val Glu Ile
 100 105 110

Ser Arg Phe Trp Val Leu Glu Ile Ile Arg Trp Leu Glu Lys Pro Ser
 115 120 125

Thr Ser Arg
 130

<210> 417

<211> 137

<212> PRT

<213> Streptococcus pneumoniae

<400> 417

Met Ser Lys Leu Leu Asp Lys Ile Leu Ser Arg Glu Asn Met Leu Glu
 1 5 10 15

Ala Tyr Asn Gln Val Lys Ser Asn Lys Gly Ser Ala Gly Ile Asp Gly
 20 25 30

Met Thr Ile Glu Glu Met Asp Asn Tyr Leu Arg Gln Asn Trp Arg Leu
 35 40 45

Thr Lys Glu Leu Ile Lys Gln Arg Lys Tyr Lys Pro Gln Pro Val Leu
 50 55 60

Arg Val Glu Ile Pro Lys Pro Asp Gly Gly Ile Arg Gln Leu Gly Ile
 65 70 75 80

Pro Thr Val Met Asp Arg Met Ile Gln Gln Ala Ile Val Gln Val Met
85 90 95

Ser Pro Ile Cys Glu Pro His Phe Ser Asp Thr Ser Tyr Gly Phe Arg
100 105 110

Pro Asn Arg Ser Cys Glu Lys Ala Ile Met Lys Leu Leu Glu Tyr Leu
115 120 125

Asn Asp Gly Tyr Glu Trp Ile Val Asp
130 135

<210> 418

<211> 164

<212> PRT

<213> Streptococcus pneumoniae

<400> 418

Met Arg Gln Leu Lys Arg Val Gly Val Phe Leu Leu Leu Pro Phe Phe
1 5 10 15

Val Leu Ile Asp Ala His Ile Ser Gln Leu Leu Gly Ser Phe Phe Pro
20 25 30

His Val His Leu Ala Ser His Phe Leu Phe Leu Phe Leu Phe Glu
35 40 45

Thr Ile Glu Val Ser Glu Tyr Leu Tyr Leu Val Tyr Cys Phe Val Ile
50 55 60

Gly Leu Val Tyr Asp Val Tyr Phe Phe His Leu Ile Gly Ile Thr Thr
65 70 75 80

Leu Leu Phe Ile Leu Leu Gly Ala Phe Leu His Lys Leu Asn Ser Val
85 90 95

Ile Leu Leu Asn Arg Trp Thr Arg Met Leu Ala Met Ile Val Leu Thr
100 105 110

Phe Leu Phe Glu Met Gly Ser Tyr Leu Leu Ala Phe Met Val Gly Leu
115 120 125

Thr Val Asp Ser Met Ser Ile Phe Ile Val Tyr Ser Leu Val Pro Thr
 130 135 140

Met Ile Leu Asn Phe Leu Trp Ile Thr Val Phe Gln Phe Ile Phe Glu
 145 150 155 160

Lys Tyr Tyr Leu

<210> 419
 <211> 427
 <212> PRT
 <213> Streptococcus pneumoniae

<400> 419

Met Thr Lys Val Val Phe Glu Glu Lys Tyr Tyr Pro Ala Val Lys Glu
 1 5 10 15

Met Val Tyr Arg Thr Arg Leu Ala Asn Gly Leu Thr Val Ala Leu Leu
 20 25 30

Pro Lys Lys Glu Phe Lys Glu Val Tyr Gly Ser Val Thr Val Gln Phe
 35 40 45

Gly Ser Val Asp Thr Phe Val Thr Glu Val Asp Gly Asp Val Lys Gln
 50 55 60

Tyr Pro Gly Gly Ile Ala His Phe Leu Glu His Lys Leu Phe Glu Arg
 65 70 75 80

Glu Asp Ser Ser Asp Leu Met Ser Ala Phe Thr Ser Leu Gly Ala Asp
 85 90 95

Ser Asn Ala Phe Thr Ser Phe Thr Lys Thr Asn Tyr Leu Phe Ser Ala
 100 105 110

Thr Asp Tyr Phe Leu Glu Asn Leu Asp Leu Leu Asp Glu Leu Val Thr
 115 120 125

Ser Ala His Phe Thr Glu Ala Ser Ile Leu Thr Glu Gln Asp Ile Ile
 130 135 140

Gln Gln Glu Arg Glu Met Tyr Gln Asp Asp Pro Asp Ser Cys Leu Phe
 145 150 155 160

Phe Ser Thr Leu Ala Asn Leu Tyr Pro Gly Thr Pro Leu Ala Thr Asp
 165 170 175

Ile Val Gly Ser Glu Glu Ser Ile Ser Gln Ile Asn Leu Thr Asn Leu
 180 185 190

Gln Glu Asn Phe Thr Lys Phe Tyr Lys Pro Val Asn Met Ser Leu Phe
 195 200 205

Leu Val Gly Asn Phe Asp Val Glu Arg Val Gln Asp Tyr Phe Glu Ser
 210 215 220

Lys Glu Leu Lys Asp Ser Asp Phe Gln Glu Val Ala Arg Glu Lys Leu
 225 230 235 240

Phe Leu Gln Pro Val Lys Pro Thr Asp Ser Met Arg Met Glu Val Ser
 245 250 255

Ser Pro Lys Leu Ala Ile Gly Val Arg Gly Lys Arg Glu Val Ser Glu
 260 265 270

Ala Asp Cys Tyr Arg His His Ile Leu Leu Lys Leu Leu Phe Ala Met
 275 280 285

Met Phe Gly Trp Thr Ser Asp Arg Phe Gln Lys Cys Tyr Glu Ser Gly
 290 295 300

Lys Ile Asp Ala Ser Leu Ser Leu Glu Val Glu Ile Thr Ser Arg Phe
 305 310 315 320

His Phe Val Met Leu Thr Met Asp Thr Lys Glu Pro Val Ala Leu Ser
 325 330 335

His Gln Phe Arg Lys Ala Ile Arg Asn Phe Thr Lys Asp Leu Asp Ile
 340 345 350

Thr Glu Glu His Leu Asp Ile Ile Lys Arg Glu Met Phe Gly Glu Phe
 355 360 365

Phe Ser Ser Met Asn Ser Leu Glu Phe Ile Ala Thr Gln Tyr Asp Ala
 370 375 380

Phe Glu Asn Gly Glu Ile Ile Phe Asp Leu Pro Lys Ile Leu Gln Glu
 385 390 395 400

Ile Thr Leu Glu Asp Val Leu Asp Ala Gly His His Leu Ile Asp Asp
 405 410 415

Gly Asp Ile Val Asp Phe Thr Ile Phe Pro Ser
 420 425

<210> 420

<211> 540

<212> PRT

<213> Streptococcus pneumoniae

<400> 420

Met Leu Thr Val Ser Asp Val Ser Leu Arg Phe Ser Asp Arg Lys Leu
 1 5 10 15

Phe Asp Asp Val Asn Ile Lys Phe Thr Glu Gly Asn Thr Tyr Gly Leu
 20 25 30

Ile Gly Ala Asn Gly Ala Gly Lys Ser Thr Phe Leu Lys Ile Leu Ala
 35 40 45

Gly Asp Ile Glu Pro Thr Thr Gly His Ile Ser Leu Gly Pro Asp Glu
 50 55 60

Arg Leu Ser Val Leu Arg Gln Asn His Phe Asp Tyr Glu Asp Glu Arg
 65 70 75 80

Ala Ile Asp Val Val Ile Met Gly Asn Glu Lys Leu Tyr Ser Ile Met
 85 90 95

Lys Glu Lys Asp Ala Ile Tyr Met Lys Glu Asp Phe Ser Asp Glu Asp
 100 105 110

Gly Val Arg Ala Ala Glu Leu Glu Gly Glu Phe Ala Glu Leu Gly Gly
 115 120 125

Trp Glu Ala Glu Ser Glu Ala Ser Gln Leu Leu Gln Asn Leu Asn Ile
 130 135 140

Pro Glu Glu Leu His Tyr Gln Asn Met Ser Glu Leu Ala Asn Gly Glu
 145 150 155 160

Lys Val Lys Val Leu Leu Ala Lys Ala Leu Phe Gly Lys Pro Asp Val
 165 170 175

Leu Leu Leu Asp Glu Pro Thr Asn Gly Leu Asp Ile Gln Ser Ile Thr
 180 185 190

Trp Leu Glu Asp Phe Leu Ile Asp Phe Asp Asn Thr Val Ile Val Val
 195 200 205

Ser His Asp Arg His Phe Leu Asn Lys Val Cys Thr His Met Ala Asp
 210 215 220

Leu Asp Phe Gly Lys Ile Lys Leu Tyr Val Gly Asn Tyr Asp Phe Trp
 225 230 235 240

Lys Glu Ser Ser Glu Leu Ala Ala Lys Leu Leu Ala Asp Arg Asn Ala
 245 250 255

Lys Ala Glu Glu Lys Ile Lys Gln Leu Gln Glu Phe Val Ala Arg Phe
 260 265 270

Ser Ala Asn Ala Ser Lys Ser Arg Gln Ala Thr Ser Arg Lys Lys Met
 275 280 285

Leu Asp Lys Ile Glu Leu Glu Glu Ile Val Pro Ser Ser Arg Lys Tyr
 290 295 300

Pro Phe Ile Asn Phe Lys Ala Glu Arg Glu Ile Gly Asn Asp Leu Leu
 305 310 315 320

Thr Val Glu Asn Leu Thr Val Lys Ile Asp Gly Glu Thr Ile Leu Asp
 325 330 335

Asn Ile Ser Phe Ile Leu Arg Pro Asp Asp Lys Thr Ala Leu Ile Gly
 340 345 350

Gln Asn Asp Ile Gln Thr Thr Ala Leu Ile Arg Ala Ile Met Gly Asp
355 360 365

Ile Asp Tyr Glu Gly Thr Val Lys Trp Gly Val Thr Thr Ser Gln Ser
370 375 380

Tyr Leu Pro Lys Asp Asn Ser Ala Asp Phe Ala Gly Gly Glu Ser Ile
385 390 395 400

Leu Asp Trp Leu Arg Gln Phe Ala Ser Lys Glu Glu Asp Asp Asn Thr
405 410 415

Phe Leu Arg Gly Phe Leu Gly Arg Met Leu Phe Ser Gly Asp Glu Val
420 425 430

Asn Lys Pro Val Asn Val Leu Ser Gly Gly Glu Lys Val Arg Val Met
435 440 445

Leu Ser Lys Leu Met Leu Leu Lys Ser Asn Val Leu Val Leu Asp Asp
450 455 460

Pro Thr Asn His Leu Asp Leu Glu Ser Ile Ser Ser Leu Asn Asp Gly
465 470 475 480

Leu Lys Asn Phe Lys Glu Ser Ile Ile Phe Ala Ser His Asp His Glu
485 490 495

Phe Ile Gln Thr Leu Ala Asn His Ile Ile Val Leu Ser Lys Asn Gly
500 505 510

Val Ile Asp Arg Ile Asp Glu Thr Tyr Asp Glu Phe Leu Glu Asn Ala
515 520 525

Glu Val Gln Ala Lys Val Lys Glu Leu Trp Lys Asp
530 535 540

<210> 421

<211> 850

<212> FRT

<213> Streptococcus pneumoniae

<400> 421

Met Lys Ser Phe Leu Lys Thr Tyr Arg Thr Tyr Phe Ile Ser Phe Ile
 1 5 10 15

Ile Pro Val Val Ile Met Ser Gly Val Tyr Leu Ser Gln Ser Ile Tyr
 20 25 30

Trp Asn Ser Asp Asn Ser Pro Leu Leu Gly Asp Gly Phe His Gln Tyr
 35 40 45

Val Ile Phe Asp Val Ala Leu Arg Asn Ile Leu His Gly Asn Ser Asn
 50 55 60

Leu Phe Tyr Thr Phe Thr Ser Gly Leu Gly Leu Asn Phe Tyr Ala Leu
 65 70 75 80

Ser Ser Tyr Tyr Leu Gly Ser Phe Leu Ala Pro Leu Val Tyr Phe Phe
 85 90 95

Asp Leu Thr Asn Met Pro Asp Ala Ile Tyr Leu Thr Thr Leu Leu Lys
 100 105 110

Phe Gly Leu Ile Gly Leu Ser Thr Phe Phe Ser Leu Asn Lys Leu Phe
 115 120 125

Gln Ser Ile Pro Gln Ile Leu Lys Leu Ala Leu Ser Thr Ser Tyr Ala
 130 135 140

Leu Met Ser Phe Thr Val Ser Gln Leu Glu Ile Lys Thr Trp Leu Asp
 145 150 155 160

Val Phe Ile Leu Ile Pro Leu Ile Ile Thr Gly Leu His Leu Leu Ile
 165 170 175

Thr Glu Lys Lys Leu Leu Leu Tyr Phe Thr Ser Leu Ser Ile Leu Phe
 180 185 190

Ile Gln Asn Tyr Tyr Phe Gly Tyr Met Thr Val Leu Phe Leu Ile Phe
 195 200 205

Trp Tyr Leu Cys Gln Ile Ser Trp Asp Phe Lys Thr Arg Lys Ser Ser

210	215	220
Val Leu Asp Phe Ile Val	Ile Ser Phe Leu Ala Gly Met Ala Ser Leu	
225	230	235 240
Ile Met Thr Leu Pro Thr	Leu Phe Asp Leu Gln Thr His Gly Glu Lys	
	245	250 255
Leu Thr Glu Val Thr Lys Phe Gln Thr Glu Ser Ser Trp Tyr Leu Asp		
	260	265 270
Leu Phe Ala Lys Gln Phe Ile Gly Ser Phe Asp Thr Thr Lys Tyr Gly		
	275	280 285
Ala Ile Pro Met Ile Phe Val Gly Leu Phe Pro Phe Ile Leu Thr Ile		
	290	295 300
Leu Phe Phe Thr Leu Lys Ser Ile Lys Phe His Val Lys Leu Ile Tyr		
305	310	315 320
Val Ile Phe Phe Ala Phe Leu Ile Ala Ser Phe Tyr Ile Glu Ala Leu		
	325	330 335
Asp Leu Phe Trp Gln Gly Met His Thr Pro Asn Met Phe Leu His Arg		
	340	345 350
Tyr Ala Trp Ile Phe Ser Thr Leu Leu Ile Tyr Thr Ala Ala Glu Val		
	355	360 365
Leu Lys Arg Leu Lys Glu Leu Lys Val Trp Asn Phe Leu Val Ser Leu		
	370	375 380
Phe Leu Val Val Ala Gly Phe Leu Ala Thr Ile Tyr Leu Lys Ser His		
385	390	395 400
Tyr Ser Phe Leu Thr Asp Leu Asn Ile Leu Leu Thr Leu Glu Phe Leu		
	405	410 415
Val Val Tyr Ser Leu Leu Leu Leu Ala Val Ile Lys Lys Phe Ile Ser		
	420	425 430

Val Asn Leu Phe Ala Ile Leu Ile Ser Leu Phe Ile Leu Val Glu Met
 435 440 445

Ser Leu Asn Ala Ser Ser Gln Met Asp Gly Ile Ala Lys Glu Trp Gly
 450 455 460

Phe Ala Ser Arg Ser Ala Tyr Ser Arg Asp Ile Pro Ala Met Glu Ser
 465 470 475 480

Phe Ser Thr Tyr Ile Gly Asn Gln Phe Thr Arg Thr Glu Lys Leu Gln
 485 490 495

Thr Gln Thr Gly Asn Asp Ser Met Lys Phe Asn Tyr Asn Gly Ile Ser
 500 505 510

Gln Phe Ser Ser Val Arg Asn Arg Ser Ser Ser Ser Thr Leu Asp Lys
 515 520 525

Leu Gly Phe Lys Ser Ser Gly Thr Asn Leu Asn Leu Arg Tyr Ala Asn
 530 535 540

Asn Ser Ile Leu Ala Asp Ser Leu Phe Gly Ile Gln Tyr Asn Ile Ser
 545 550 555 560

Asp Ser Pro Ile Asp Lys Tyr Gly Phe Lys Asp Ile Tyr Gln Lys Asp
 565 570 575

Asn Leu Thr Leu Tyr Glu Asn Gln Tyr Ser Leu Pro Ile Ala Val Ala
 580 585 590

Ser Gln Ser Val Tyr Asn Asp Val Lys Phe Asn Glu His Thr Leu Asp
 595 600 605

Asn Gln Ala Ser Phe Leu Asn Gln Leu Ala Asn Val Asn Phe Asp Tyr
 610 615 620

Phe Ser Pro Ile Pro Tyr Glu Lys Thr Glu Lys Ile Glu Asn Thr Asn
 625 630 635 640

Asp Leu Ile Ser Val Thr Ser Ser Ser Asn Glu Asp Ala Ala Ile Gln
 645 650 655

Tyr Gln Ile Glu Val Pro Glu Asn Ser Gln Val Tyr Leu Ser Phe Ile
 660 665 670
 Asn Leu His Phe Ser Asn Asp Lys Gln Lys Lys Val Asp Ile Leu Val
 675 680 685
 Asn Gly Glu Lys Lys Thr Phe Thr Thr Asp Asn Val Phe Ser Phe Phe
 690 695 700
 Asn Leu Gly Tyr Thr Lys Glu Lys Lys Thr Phe Asn Ile Asn Val Ser
 705 710 715 720
 Phe Pro Gly Asn Ser Gln Val Ser Phe Glu Ser Pro Thr Phe Tyr Arg
 725 730 735
 Leu Asp Thr Lys Thr Phe Thr Glu Ala Ile Gln Lys Ile Lys Glu Gln
 740 745 750
 Pro Val Thr Val Ser Thr Ser Lys Asn Lys Val Phe Ala Thr Tyr Asp
 755 760 765
 Val Gln Gln Asp Thr Ser Ile Phe Phe Thr Ile Pro Tyr Asp Lys Gly
 770 775 780
 Trp Ser Ala Tyr Gln Asp Gly Lys Lys Ile Glu Ile Lys Gln Ala Gln
 785 790 795 800
 Thr Gly Phe Met Lys Val Asp Ile Pro Lys Gly Lys Gly Thr Ile Thr
 805 810 815
 Leu Ser Phe Ile Pro Asn Gly Phe Ile Thr Gly Ala Ile Cys Ser Phe
 820 825 830
 Thr Ser Leu Leu Leu Phe Gly Ile Tyr Asn His Arg Arg Lys Ser Ser
 835 840 845
 Lys Ala
 850

<210> 422

<211> 250

<212> PRT

<213> Streptococcus pneumoniae

<400> 422

Met Lys Val Leu Ile Leu Glu Asp Val Ile Glu His Gln Val Arg Leu
 1 5 10 15

Glu Arg Ile Leu Asp Glu Ile Ser Lys Glu Ser Asn Ile Pro Ile Ser
 20 25 30

Tyr Lys Thr Thr Gly Lys Val Arg Glu Phe Glu Glu Tyr Ile Glu Asn
 35 40 45

Asp Glu Val Asn Gln Leu Tyr Phe Leu Asp Ile Asp Ile His Gly Ile
 50 55 60

Glu Lys Lys Gly Phe Glu Val Ala Gln Leu Ile Arg His Tyr Asn Pro
 65 70 75 80

Tyr Ala Ile Ile Val Phe Ile Thr Ser Arg Ser Glu Phe Ala Thr Leu
 85 90 95

Thr Tyr Lys Tyr Gln Val Ser Ala Leu Asp Phe Val Asp Lys Asp Ile
 100 105 110

Asn Asp Glu Met Phe Lys Lys Arg Ile Glu Gln Asn Ile Phe Tyr Thr
 115 120 125

Lys Ser Met Leu Leu Glu Asn Glu Asp Val Val Asp Tyr Phe Asp Tyr
 130 135 140

Asn Tyr Lys Gly Asn Asp Leu Lys Ile Pro Tyr His Asp Ile Leu Tyr
 145 150 155 160

Ile Glu Thr Thr Gly Val Ser His Lys Leu Arg Ile Ile Gly Lys Asn
 165 170 175

Phe Ala Lys Glu Phe Tyr Gly Thr Met Thr Asp Ile Gln Glu Lys Asp
 180 185 190

Lys His Thr Gln Arg Phe Tyr Ser Pro His Lys Ser Phe Leu Val Asn
 195 200 205

Ile Gly Asn Ile Arg Glu Ile Asp Arg Lys Asn Leu Glu Ile Val Phe
 210 215 220

Tyr Glu Asp His Arg Cys Pro Ile Ser Arg Leu Lys Ile Arg Lys Leu
 225 230 235 240

Lys Asp Ile Leu Glu Lys Lys Ser Gln Lys
 245 250

<210> 423
 <211> 441
 <212> PRT
 <213> Streptococcus pneumoniae

<400> 423

Met Asp Leu Leu Gly Phe Gly Thr Val Ile Val His Phe Leu Ile Ile
 1 5 10 15

Ser His Ser Tyr Arg Leu Ile Cys Lys Gly Arg Ile Asn Arg Lys Glu
 20 25 30

Leu Tyr Val Phe Gly Ala Tyr Thr Leu Leu Thr Glu Ile Val Leu Glu
 35 40 45

Phe Ser Phe Tyr Leu Leu Tyr Leu Asp Lys Ile Gly Ile Glu Arg Phe
 50 55 60

Leu Phe Pro Leu Gly Leu Tyr Ser Tyr Phe Arg Trp Met Lys Gln Tyr
 65 70 75 80

Glu Arg Asp Arg Gly Leu Phe Leu Ser Leu Leu Leu Ser Leu Tyr
 85 90 95

Glu Ser Thr His Asn Phe Leu Ser Val Ile Phe Ser Ser Ile Thr Gly
 100 105 110

Asp Asn Phe Val Leu Gln Tyr His Phe Pro Phe Phe Phe Val Val Thr
 115 120 125

Val Leu Thr Tyr Phe Val Thr Leu Lys Ile Ile Tyr Tyr Phe His Leu
 130 135 140

Glu Leu Ala Tyr Phe Asp Lys Asp Tyr Leu Tyr Pro Phe Leu Lys Lys
 145 150 155 160
 Val Phe Phe Ala Leu Leu Leu Leu His Ile Val Ser Phe Val Ser Asp
 165 170 175
 Met Val Ser Thr Ile Lys His Leu Asn Ser Phe Gly Ser Ile Leu Ser
 180 185 190
 Ser Ile Val Phe Ile Ser Leu Leu Leu Thr Phe Phe Ala Met Asn Ser
 195 200 205
 His Lys Val Gln Met Glu Lys Glu Ile Ala Leu Lys Gln Lys Lys Phe
 210 215 220
 Glu Gln Lys His Leu Gln Asn Tyr Thr Asp Glu Ile Val Gly Leu Tyr
 225 230 235 240
 Asn Glu Ile Arg Gly Phe Arg His Asp Tyr Ala Gly Met Leu Val Ser
 245 250 255
 Met Gln Met Ala Ile Asp Ser Gly Asn Leu Gln Glu Ile Asp Arg Ile
 260 265 270
 Tyr Asn Glu Val Leu Val Lys Ala Asn His Lys Leu Arg Ser Asp Lys
 275 280 285
 Tyr Thr Tyr Phe Asp Leu Asn Asn Ile Glu Asp Ser Ala Leu Arg Ser
 290 295 300
 Leu Val Ala Gln Ser Ile Val Tyr Ala Arg Asn Asn Gly Val Glu Phe
 305 310 315 320
 Thr Leu Glu Val Lys Asp Thr Ile Thr Lys Leu Pro Ile Glu Leu Leu
 325 330 335
 Asp Leu Val Arg Ile Met Ser Val Leu Leu Asn Asn Ala Val Glu Gly
 340 345 350
 Ser Ala Asp Ser Tyr Lys Lys Gln Met Glu Val Ala Val Ile Lys Met

355 360 365
 Glu Thr Glu Thr Val Ile Val Ile Gln Asn Ser Cys Lys Met Thr Met
 370 375 380
 Thr Pro Ser Gly Asp Leu Phe Ala Leu Gly Phe Ser Thr Lys Gly Arg
 385 390 395 400
 Asn Arg Gly Val Gly Leu Asn Asn Val Lys Glu Leu Leu Asp Lys Tyr
 405 410 415
 Asn Asn Ile Ile Leu Glu Thr Glu Met Glu Gly Ser Thr Phe Arg Gln
 420 425 430
 Ile Ile Arg Phe Lys Arg Glu Phe Glu
 435 440
 <210> 424
 <211> 122
 <212> PRT
 <213> Streptococcus pneumoniae
 <400> 424
 Met Ser Lys Asn Ile Val Gln Leu Asn Asn Ser Phe Ile Gln Asn Glu
 1 5 10 15
 Tyr Gln Arg Arg Arg Tyr Leu Met Lys Glu Arg Gln Lys Arg Asn Arg
 20 25 30
 Phe Met Gly Gly Val Leu Ile Leu Ile Met Leu Leu Phe Ile Leu Pro
 35 40 45
 Thr Phe Asn Leu Ala Gln Ser Tyr Gln Gln Leu Leu Gln Arg Arg Gln
 50 55 60
 Gln Leu Ala Asp Leu Gln Thr Gln Tyr Gln Thr Leu Ser Asp Glu Lys
 65 70 75 80
 Asp Lys Glu Thr Ala Phe Ala Thr Lys Leu Lys Asp Glu Asp Tyr Ala
 85 90 95
 Ala Lys Tyr Thr Arg Ala Lys Tyr Tyr Tyr Ser Lys Ser Arg Glu Lys

100 105 110
 Val Tyr Thr Ile Pro Asp Leu Leu Gln Arg
 115 120

 <210> 425
 <211> 420
 <212> PRT
 <213> Streptococcus pneumoniae

 <400> 425
 Met Glu Glu Val Glu Val Ala Glu Val Lys Asn Ala Arg Val Ser Leu
 1 5 10 15

 Thr Gly Glu Lys Thr Lys Pro Met Lys Leu Ala Glu Val Thr Ser Ile
 20 25 30

 Asn Val Asn Arg Thr Lys Thr Glu Met Glu Glu Phe Asn Arg Val Leu
 35 40 45

 Gly Gly Gly Val Val Pro Gly Ser Leu Val Leu Ile Gly Gly Asp Pro
 50 55 60

 Gly Ile Gly Lys Ser Thr Leu Leu Leu Gln Val Ser Thr Gln Leu Ser
 65 70 75 80

 Gln Val Gly Thr Val Leu Tyr Val Ser Gly Glu Glu Ser Ala Gln Gln
 85 90 95

 Ile Lys Leu Arg Ala Glu Arg Leu Gly Asp Ile Asp Ser Glu Phe Tyr
 100 105 110

 Leu Tyr Ala Glu Thr Asn Met Gln Ser Val Arg Ala Glu Val Glu Arg
 115 120 125

 Ile Gln Pro Asp Phe Leu Ile Ile Asp Ser Ile Gln Thr Ile Met Ser
 130 135 140

 Pro Glu Ile Ser Gly Val Gln Gly Ser Val Ser Gln Val Arg Glu Val
 145 150 155 160

 Thr Ala Glu Leu Met Gln Leu Ala Lys Thr Asn Asn Ile Ala Ile Phe

	165		170		175
Ile Val Gly	His Val Thr Lys Glu Gly Thr Leu Ala Gly Pro Arg Met				
	180		185		190
Leu Glu His Met Val Asp Thr Val Leu Tyr Phe Glu Gly Glu Arg His					
	195		200		205
His Thr Phe Arg Ile Leu Arg Ala Val Lys Asn Arg Phe Gly Ser Thr					
	210		215		220
Asn Glu Ile Gly Ile Phe Glu Met Gln Ser Gly Gly Leu Val Glu Val					
	225		230		235
Leu Asn Pro Ser Gln Val Phe Leu Glu Glu Arg Leu Asp Gly Ala Thr					
			245		250
Gly Ser Ser Ile Val Val Thr Met Glu Gly Thr Arg Pro Ile Leu Ala					
			260		265
Glu Val Gln Ala Leu Val Thr Pro Thr Met Phe Gly Asn Ala Lys Arg					
			275		280
Thr Thr Thr Gly Leu Asp Phe Asn Arg Ala Ser Leu Ile Met Ala Val					
			290		295
Leu Glu Lys Arg Ala Gly Leu Leu Leu Gln Asn Gln Asp Ala Tyr Leu					
			305		310
Lys Ser Ala Gly Gly Val Lys Leu Asp Glu Pro Ala Ile Asp Leu Ala					
			325		330
Val Ala Val Ala Ile Ala Ser Ser Tyr Lys Asp Lys Pro Thr Asn Pro					
			340		345
Gln Glu Cys Phe Val Gly Glu Leu Gly Leu Thr Gly Glu Ile Arg Arg					
			355		360
Val Asn Arg Ile Glu Gln Arg Ile Asn Glu Ala Ala Lys Leu Gly Phe					
			370		375

Thr Lys Ile Tyr Val Pro Lys Asn Ser Leu Thr Gly Ile Thr Leu Pro
 385 390 395 400

Lys Glu Ile Gln Val Ile Gly Val Thr Thr Ile Gln Glu Val Leu Lys
 405 410 415

Lys Val Phe Ala
 420

<210> 426

<211> 207

<212> PRT

<213> Streptococcus pneumoniae

<400> 426

Met Pro Lys Lys Val Leu Ile Leu Ser Leu Leu Gly Gly Met Phe Leu
 1 5 10 15

Ser Gly Trp Leu Ser Ser Phe Ala Asn Thr Tyr Ile His Asp Leu Leu
 20 25 30

Gly Val Leu Phe Pro Asp Ser Pro Phe Leu Asn Ala Phe Glu Ser Ala
 35 40 45

Ile Ala Ala Pro Leu Val Glu Glu Pro Leu Lys Leu Leu Ser Leu Val
 50 55 60

Phe Val Leu Ala Leu Ile Pro Val Arg Lys Leu Lys Ser Leu Phe Leu
 65 70 75 80

Leu Gly Ile Ala Ser Gly Leu Gly Phe Gln Met Ile Lys Asp Ile Gly
 85 90 95

Tyr Ile Arg Thr Asp Leu Pro Glu Gly Phe Asp Phe Thr Ile Ser Arg
 100 105 110

Ile Leu Glu Arg Ile Ile Ser Gly Ile Ala Ser His Trp Thr Phe Ser
 115 120 125

Gly Leu Ala Val Val Gly Val Tyr Leu Leu Tyr Arg Ala Tyr Lys Gly
 130 135 140

Gln Lys Val Gly Lys Lys Gln Gly Leu Ile Phe Leu Gly Leu Ala Leu
 145 150 155 160

Gly Thr His Phe Leu Phe Asn Ser Pro Phe Val Glu Leu Glu Thr Glu
 165 170 175

Leu Pro Leu Ala Ile Pro Val Val Thr Ala Ile Ala Leu Tyr Gly Phe
 180 185 190

Tyr His Ala Tyr Cys Phe Val Glu Lys His Asn Glu Leu Met Thr
 195 200 205

<210> 427

<211> 717

<212> PRT

<213> Streptococcus pneumoniae

<400> 427

Met Lys Phe Gly Lys Arg His Tyr Arg Pro Gln Val Asp Gln Met Asp
 1 5 10 15

Cys Gly Val Ala Ser Leu Ala Met Val Phe Gly Tyr Tyr Gly Ser Tyr
 20 25 30

Tyr Phe Leu Ala His Leu Arg Glu Leu Ala Lys Thr Thr Met Asp Gly
 35 40 45

Thr Thr Ala Leu Gly Leu Val Lys Val Ala Glu Glu Ile Gly Phe Glu
 50 55 60

Thr Arg Ala Ile Lys Ala Asp Met Thr Leu Phe Asp Leu Pro Asp Leu
 65 70 75 80

Thr Phe Pro Phe Val Ala His Val Leu Lys Glu Gly Lys Leu Leu His
 85 90 95

Tyr Tyr Val Val Thr Gly Gln Asp Lys Asp Ser Ile His Ile Ala Asp
 100 105 110

Pro Asp Pro Gly Val Lys Leu Thr Lys Leu Pro Arg Glu Arg Phe Glu
 115 120 125

Glu Glu Trp Thr Gly Val Thr Leu Phe Met Ala Pro Ser Pro Asp Tyr
 130 135 140

Lys Pro His Lys Glu Gln Lys Asn Gly Leu Leu Ser Phe Ile Pro Ile
 145 150 155 160

Leu Val Lys Gln Arg Gly Leu Ile Ala Asn Ile Val Leu Ala Thr Leu
 165 170 175

Leu Val Thr Val Ile Asn Ile Val Gly Ser Tyr Tyr Leu Gln Ser Ile
 180 185 190

Ile Asp Thr Tyr Val Pro Asp Gln Met Arg Ser Thr Leu Gly Ile Ile
 195 200 205

Ser Ile Gly Leu Val Ile Val Tyr Ile Phe Gln Gln Ile Leu Ser Tyr
 210 215 220

Ala Gln Glu Tyr Leu Leu Leu Val Leu Gly Gln Arg Leu Ser Ile Asp
 225 230 235 240

Val Ile Leu Ser Tyr Ile Lys His Val Phe His Leu Pro Met Ser Phe
 245 250 255

Phe Ala Thr Arg Arg Thr Gly Glu Ile Val Ser Arg Phe Thr Asp Ala
 260 265 270

Asn Ser Ile Ile Asp Ala Leu Ala Ser Thr Ile Leu Ser Ile Phe Leu
 275 280 285

Asp Val Ser Thr Val Val Ile Ile Ser Leu Val Leu Phe Ser Gln Asn
 290 295 300

Thr Asn Leu Phe Phe Met Thr Leu Leu Ala Leu Pro Ile Tyr Thr Val
 305 310 315 320

Ile Ile Phe Ala Phe Met Lys Pro Phe Glu Lys Met Asn Arg Asp Thr
 325 330 335

Met Glu Ala Asn Ala Val Leu Ser Ser Ser Ile Ile Glu Asp Ile Asn
 340 345 350

Gly Ile Glu Thr Ile Lys Ser Leu Thr Ser Glu Ser Gln Arg Tyr Gln
 355 360 365
 Lys Ile Asp Lys Glu Phe Val Asp Tyr Leu Lys Lys Ser Phe Thr Tyr
 370 375 380
 Ser Arg Ala Glu Ser Gln Gln Lys Ala Leu Lys Lys Val Ala His Leu
 385 390 395 400
 Leu Leu Asn Val Gly Ile Leu Trp Met Gly Ala Val Leu Val Met Asp
 405 410 415
 Gly Lys Met Ser Leu Gly Gln Leu Ile Thr Tyr Asn Thr Leu Leu Val
 420 425 430
 Tyr Phe Thr Asn Pro Leu Glu Asn Ile Ile Asn Leu Gln Thr Lys Leu
 435 440 445
 Gln Thr Ala Gln Val Ala Asn Asn Arg Leu Asn Glu Val Tyr Leu Val
 450 455 460
 Ala Ser Glu Phe Glu Glu Lys Lys Thr Val Glu Asp Leu Ser Leu Met
 465 470 475 480
 Lys Gly Asp Met Thr Phe Lys Gln Val His Tyr Lys Tyr Gly Tyr Gly
 485 490 495
 Arg Asp Val Leu Ser Asp Ile Asn Leu Thr Val Pro Gln Gly Ser Lys
 500 505 510
 Val Ala Phe Val Gly Ile Ser Gly Ser Gly Lys Thr Thr Leu Ala Lys
 515 520 525
 Met Met Val Asn Phe Tyr Asp Pro Ser Gln Gly Glu Ile Ser Leu Gly
 530 535 540
 Ser Val Asn Leu Asn Gln Ile Asp Lys Lys Ala Leu Arg Gln Tyr Ile
 545 550 555 560
 Asn Tyr Leu Ser Gln Gln Pro Tyr Val Phe Asn Gly Thr Ile Leu Glu
 565 570 575

Asn Leu Leu Leu Gly Ala Lys Glu Gly Thr Thr Gln Glu Asp Ile Leu
580 585 590

Arg Ala Val Glu Leu Ala Glu Ile Arg Glu Asp Ile Glu Arg Met Pro
595 600 605

Leu Asn Tyr Gln Thr Glu Leu Thr Ser Asp Gly Ala Gly Ile Ser Gly
610 615 620

Gly Gln Arg Gln Arg Ile Ala Leu Ala Arg Ala Leu Leu Thr Asp Ala
625 630 635 640

Pro Val Leu Ile Leu Asp Glu Ala Thr Ser Ser Leu Asp Ile Leu Thr
645 650 655

Glu Lys Arg Ile Val Asp Asn Leu Ile Ala Leu Asp Lys Thr Leu Ile
660 665 670

Phe Ile Ala His Arg Leu Thr Ile Ala Glu Arg Thr Glu Lys Val Val
675 680 685

Val Leu Asp Gln Gly Lys Ile Val Glu Glu Gly Lys His Ala Asp Leu
690 695 700

Leu Ala Gln Gly Gly Phe Tyr Ala His Leu Val Asn Ser
705 710 715

<210> 428

<211> 319

<212> PRT

<213> Streptococcus pneumoniae

<400> 428

Met Asp Ile Lys Ile Lys Arg Glu Glu Ile Met Lys Lys Phe Ser Lys
1 5 10 15

Thr Leu Arg Asp Asn Trp Ile Phe Leu Leu Met Val Leu Pro Gly Ala
20 25 30

Leu Trp Leu Ile Leu Phe Phe Tyr Ile Pro Val Phe Gly Asn Val Val
35 40 45

Ala Phe Lys Asp Tyr His Met Thr Ser Asn Gly Phe Ile Asp Ser Ile
 50 55 60
 Ile Asn Ser Lys Trp Val Gly Leu Asp Asn Phe Arg Phe Leu Phe Ser
 65 70 75 80
 Ser Arg Asp Ala Phe Ile Ile Thr Arg Asn Thr Val Leu Tyr Asn Leu
 85 90 95
 Gly Phe Ile Phe Leu Gly Leu Val Val Ser Val Gly Ile Ala Ile Ile
 100 105 110
 Leu Ser Glu Leu Arg Ser Lys Arg Met Val Lys Ile Phe Gln Thr Ser
 115 120 125
 Met Leu Phe Pro Tyr Phe Leu Ser Trp Val Ile Ile Ser Phe Phe Thr
 130 135 140
 Asp Ala Phe Leu Asn Ile Asp Lys Gly Val Phe Asn His Leu Leu Glu
 145 150 155 160
 Ser Leu Gly Leu Lys Glu Val Asn Phe Tyr Ala Asp Leu Gly Ile Trp
 165 170 175
 Pro Tyr Leu Leu Leu Phe Leu Gly Ile Trp Lys Gly Phe Gly Tyr Ser
 180 185 190
 Ser Val Met Tyr Tyr Ala Thr Ile Met Gly Ile Asp Pro Thr Tyr Tyr
 195 200 205
 Glu Ala Ala Thr Val Asp Gly Ala Ser Lys Trp Gln Arg Ile Arg Asn
 210 215 220
 Val Thr Ile Pro Gln Leu Thr Pro Leu Val Thr Val Leu Thr Ile Leu
 225 230 235 240
 Ala Val Gly Asn Ile Phe Arg Ala Asp Phe Gly Leu Phe Tyr Gln Ile
 245 250 255
 Pro His Asn Ala Gly Gln Leu Tyr Asn Val Thr Asn Val Leu Asp Val

260 265 270
 Tyr Val Phe Asn Gly Leu Thr Gln Thr Ala Asp Ile Gly Met Ala Ala
 275 280 285
 Ala Ala Gly Leu Tyr Gln Ser Val Val Gly Leu Ile Leu Val Ile Leu
 290 295 300
 Ser Asn Leu Leu Ala Arg Arg Val Asp Pro Asn Ser Ala Leu Phe
 305 310 315
 <210> 429
 <211> 388
 <212> PRT
 <213> Streptococcus pneumoniae
 <400> 429
 Met Lys Lys Gln Ser Leu Phe Phe Val Pro Gly Ile Ile Leu Ile Gly
 1 5 10 15
 Val Ser Leu Arg Thr Pro Phe Thr Val Leu Pro Ile Ile Leu Gly Asn
 20 25 30
 Ile Ser Gln Gly Leu Glu Val Glu Val Ser Ser Leu Gly Val Leu Thr
 35 40 45
 Ser Leu Pro Leu Leu Met Phe Thr Leu Phe Ser Pro Phe Ser Thr Gln
 50 55 60
 Leu Ala Gln Lys Ile Gly Leu Glu His Leu Phe Thr Tyr Ser Leu Phe
 65 70 75 80
 Phe Leu Thr Ile Gly Ser Leu Ile Arg Leu Ile Asn Leu Pro Leu Leu
 85 90 95
 Tyr Leu Gly Thr Leu Met Val Gly Ala Ser Val Ala Val Ile Asn Val
 100 105 110
 Leu Leu Pro Ser Leu Ile Gln Ala Asn Gln Pro Lys Lys Ile Gly Phe
 115 120 125
 Leu Thr Thr Leu Tyr Val Thr Ser Met Gly Ile Ala Thr Ala Leu Ala

130	135	140
Ser Tyr Leu Ala Val Pro Ile Thr Gln Ala Ser Ser Trp Lys Gly Leu 145 150 155 160		
Ile Leu Leu Leu Thr Leu Leu Cys Leu Ala Thr Phe Leu Val Trp Leu 165 170 175		
Pro Asn His Arg Tyr Asn His Arg Leu Ala Pro Gln Thr Lys Gln Lys 180 185 190		
Ser Gln Ile Lys Val Met Arg Asn Lys Gln Val Trp Ala Ile Ile Ile 195 200 205		
Phe Ser Gly Phe Gln Ser Leu Ile Phe Tyr Thr Val Met Thr Trp Leu 210 215 220		
Pro Thr Met Ser Ile His Ala Gly Leu Ser Ser His Glu Ala Gly Leu 225 230 235 240		
Leu Thr Ser Ile Leu Ser Leu Ile Ser Ile Pro Phe Ser Met Thr Ile 245 250 255		
Pro Ser Leu Thr Thr Ser Leu Ser Thr Arg Asn Arg Gln Leu Met Leu 260 265 270		
Thr Leu Val Ser Leu Ala Gly Val Val Gly Ile Ser Met Leu Phe Phe 275 280 285		
Pro Ile Asn Asn Phe Ile Tyr Trp Leu Ala Ile His Leu Leu Ile Gly 290 295 300		
Thr Ala Thr Ser Ala Leu Phe Pro Tyr Leu Met Val Asn Phe Ser Leu 305 310 315 320		
Lys Thr Ser Ala Pro Glu Lys Thr Ala Gln Leu Ser Gly Leu Ser Gln 325 330 335		
Thr Gly Gly Tyr Ile Leu Ala Ala Phe Gly Pro Thr Leu Phe Gly Tyr 340 345 350		

Ser Phe Asp Leu Phe His Ser Trp Val Pro Ser Val Ala Ala Leu Leu
 355 360 365

Leu Ile Asp Ile Leu Met Thr Val Ala Leu Phe Thr Val Asp Arg Ala
 370 375 380

Asp Lys Ile Leu
 385

<210> 430
 <211> 150
 <212> PRT
 <213> Streptococcus pneumoniae

<400> 430

Met Asp Phe Leu Phe Ala Ala Gly Ala Phe Gly Leu Val Ile Ala Asn
 1 5 10 15

Asn Ala Ser Ile Ser Gly Ala Glu Gly Gly Cys Gln Ala Glu Val Gly
 20 25 30

Ser Ala Ser Ala Met Ser Ala Ala Ala Leu Thr Leu Ala Ala Gly Gly
 35 40 45

Thr Pro Tyr Gln Ala Ser Gln Ala Ile Ala Phe Val Ile Lys Asn Met
 50 55 60

Leu Gly Leu Ile Cys Asp Pro Val Ala Gly Leu Val Glu Val Pro Cys
 65 70 75 80

Val Lys Arg Asn Ala Met Gly Ala Ser Phe Ala Phe Ile Ala Ala Asp
 85 90 95

Met Ala Leu Ala Gly Ile Glu Ser Lys Ile Pro Val Asp Glu Val Ile
 100 105 110

Asp Ala Met Tyr Gln Val Gly Ala Ser Met Pro Thr Ala Phe Arg Glu
 115 120 125

Thr Ala Glu Gly Gly Leu Ala Thr Thr Pro Thr Gly Arg Arg Leu Gln
 130 135 140

Lys Glu Ile Phe Gly Glu
145 150

<210> 431
<211> 663
<212> DNA
<213> *Streptococcus pneumoniae*

<400> 431
ggaagtagtt ttatgaatat agcagtaatt ggtttggggc atgttgggct ggcctacgcg 60
ttactatttg catctaaata taaagttggt gcatatgata tagattctgt aaaaataaat 120
aatttaaaaa agggcattct tccatctaaa aatgaagagc ttatgaagtt tttttgcgag 180
aataacttaa atattacttt ttttgatata ttttctgaaa ttaaaaaata tattgattat 240
tatattattg cgcttccgac agattatgat gagaaaattg gtagttttaa tacatatgaa 300
atcgacaaaa cgggtatcgaa gattctgagg gtaaaacctt atggaaagat tattttaaag 360
tcaacagttc cgatcggttt ttcaacaaaa ttaaaaaggc tgtttgatgc aaaaaatata 420
atttttgtcc ctgaattttt gagagaaggt tggtctatat atgataattt atatccaagt 480
cgcatagttg ttggagatga gacagttgaa ggaagaaaaa ttgcagaggt gtctctttcg 540
attagtactc atagtactgc caatattaaa aatgttatgt tagtttctcc tactgaagca 600
gaagcaatta agcttttttc taacacattc ttactctccc gtgttgcttt tttaatgaac 660
tag 663

<210> 432
<211> 708
<212> DNA
<213> *Streptococcus pneumoniae*

<400> 432
aaaatgggtc taagaggagt tcctatgtct caaattgatc tacaaaaatt aactaagaaa 60
aaccaagagt ttgtccacat tgctacccaa caattcatca aagatgggaa aacagacgct 120
gaaatccaga ctatttttga ggaagtcatc ccccaaatcc ttgaggagca atctaagggt 180
acaactgcc gttccctata cggcgccacca actcattggg ctcatagctt cactgtcaaa 240
gagcagtacg aaaaagagca tccaaaagaa aatgatgacc caaaactgat gattatggac 300
tcagctcttt tcatcactag cctcttttgc cttgtcagcg ccctcacaa cttctttgcg 360
gcagaccaag ctttcggcta tggattgatt actctctat tagttggact ggttggtgga 420

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tttgcttctt acttgatgta ctactttgtt taccaatact atggaccaga tatggatcgc      480
agtcaacgtc cacctttctg gaaatctgta ctagtattcc tagcttctat gtctctttgg      540
ttgcttgtct tctttgcaac aagcttccta ccagctagcc ttaaccagct actggatcca      600
ttgccactag ctattattgg agcagccctc ctagcccttc gcttctatct caagaaacgc      660
ttgaatatcc gtatgtcaag tgcaggacca acacgctatc aagaataa                    708

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<210> 433
<211> 960
<212> DNA
<213> Streptococcus pneumoniae

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<400> 433
gaaaggataa gggatatgaa acaagttttt ctctctacaa caactgaatt taaagagatc      60
gatacgcttg aacgggttac ttggatcaat ctctgcaatc cgactcaaaa tgaatcactc      120
gaaatcgcca acaccttoga tattgatatt gctgaccttc gagcaccgct cgaatcgga      180
gaaatgtctc gtattaccat tgaagaacgag tataacctga ttatcttaga cgtgcgggtc      240
acggaggaaa gaaataacog cacctactac gtaaccatcc cgcttggtat tatcatcact      300
gaggaacca ttatcactac gtgtttggaa ccaactacgt tccttgatgt ctttatcaac      360
cgtcgattgc gtaatttcta taccttcatg cgttcacgtt ttatctttca aattctttat      420
cgcaatgcag agctttacct aacagccctt cgttcaatcg acogcaagag tgaacaaatc      480
gaaagtcaac tgcatcaatc aactcgtaat gaagaattga ttgagctcat ggaattggaa      540
aaaactatcg tctatttcaa ggcctccctc aaaacaaatg agcgcgtgat taagaattg      600
accagtccaa ccagcaatat caagaataac cttgaggacg aagacctgct tgaagacacc      660
ctgattgaaa cccaacagcg catcgagatg gcagatatatt atggaaacgt cttgcattct      720
atgacagaga cctttgcctc tatcatttct aacaaccaga acaacatcat gaaaaccttg      780
gcccttgtag ccacgtgcat gtccatccca accatgggtc tttctgccta cgggatgaac      840
tttaaggata atgaaatccc cetaaacgga gagccaaatg ccttctgggt aatcgctctt      900
atgcgccttg ctatgagtgt ctgcctcact ctctatctca tccataaaaa atggttctaa      960

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<210> 434
<211> 330
<212> DNA
<213> Streptococcus pneumoniae

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<400> 434
 atagacgaaa agggagaaaaa gatggcaaac aaaaaaatcc gtatccgttt gaaagcttac 60
 gaacaccgta cgcttgacac agcgggtgca aaaatcgtag aatcagctac togtacaggt 120
 gcacaagtgt cgggtccaat cccacttcca actgaacgta gcctctacac aatcattcgt 180
 gcgactcaca aatacaaaaga ctctcgcgaa caattttgaa tgcgtacaca caaacgtttg 240
 atcgatatcg ttaacccaac tcaaaaaaca gttgatgcct tgatgaaatt ggatcttcca 300
 agtgggtgtaa acgtagaaat caaactttaa 330

<210> 435
 <211> 645
 <212> DNA
 <213> Streptococcus pneumoniae

<400> 435
 aaaaggaact attttctcat gacaaaagga atcttaggga aaaaagtggt aatgactcaa 60
 atcttcaactg aagctggcga attgatccct gtaacagtta ttgaagcaac tccaaacgtt 120
 gttcttcaag ttaaaaactgt tgaacagac ggatacaacg ctatccaagt tggtttcgat 180
 gacaaacgcg aagtattgag caacaaacct gctaaaggac atgtagcgaa agctaacacg 240
 gctcctaagc gcttcattcg tgaattcaaa aacgttgaag gcttggaagt tgggtgctgaa 300
 attacagttg aaacattcgc agctggagac gttgttgacg taacgggtac ttctaaaggt 360
 aaaggtttcc aaggtgttat caaacgccac ggacaatcac gtggaccaat ggctcacgggt 420
 tctcgttacc accgtcgtcc aggttctatg gggcctgttg cacctaaccg cgtattcaaa 480
 ggtaaaaacc ttgcaggacg tatgggtggc gacgcggtaa caattcaaaa ccttgaagtt 540
 gtacaagtgt ttccagaaaa gaacgttata cttatcaaag gtaacgtacc aggtgctaag 600
 aaatctctta tcactatcaa atcagcagtt aaagctggta aataa 645

<210> 436
 <211> 645
 <212> DNA
 <213> Streptococcus pneumoniae

<400> 436
 agaaggggga aatcagtcac aatggcaaac gtaacattat ttgaccaaac tggtaagaa 60
 gctggccaag ttgttcttag cgatgcagta tttggtatcg aaccaaatga atcagttgtg 120
 tttgatgtaa tcatcagcca acgcgcaagc cttcgtcaag gaacacacgc tgttaaaaac 180

cgctctgcag tatcagggtg tggacgcaaa ccatggcgtc aaaaagggaac tggacgtgct	240
cgtaaggtt ctatccgctc accacaattg cgtggtggtg gtgttgctct cggaccaact	300
ccacgttcat acggtctcaa acttcacaa aaagtctgct gcctagctct taaatcagtt	360
tactctgaaa aagttgctga aaacaaattc gtacgtgtag acgctcttctc atttacagct	420
ccaaaaactg ctgaatttgc aaaagttctt gcagcattga gcacgattc taaagttctt	480
gtttaccttg aagaaggaaa tgaattcgca gctcttctg ctcgtaacct tccaaactg	540
aaagttgcaa ctgctacaac tgcaagtggt cttgacatcg caaatagcga caaacttctt	600
gtcacacaag cagctatctc taaaatcgag gaggttcttg cataa	645

<210> 437
 <211> 318
 <212> DNA
 <213> Streptococcus pneumoniae

<400> 437	
aatcgaggag gttcttgcac aatgaatttg tatgatgtta tcaaaaaacc tgcacacact	60
gaaagctcaa tggctcaact tgaagcagga aaatatgtat ttgaagtga cactcgtgca	120
cacaaacttt tgatcaagca agctgttgaa gctgcttctg aaggtgttaa agttgccaat	180
gttaacacaa tcaacgtaaa accaaaagct aaacgtgttg gacgttacac tgggtttact	240
aacaaaacta aaaaagctat catcacactt acagctgatt ctaaagcaat cgagttgttt	300
gtgctgaag ctgaataa	318

<210> 438
 <211> 846
 <212> DNA
 <213> Streptococcus pneumoniae

<400> 438	
ggaggaaata tcgtgggaat tcgtgtttat aaaccaacaa caaacggtcg cgttaatatg	60
acttcttttg atttcgtga aatcacaca agcactcctg aaaaatcatt gcttgttgca	120
ttgaagacga aggtgggtcg taacaacaac ggtcgtatca cagttcgtca ccaagtggt	180
ggacacaaac gtttctaccg tttggttgac ttcaaacgta ataaagacaa cgttgaagca	240
gttggttaaaa caatcgagta cgtccaaac cgttctgcaa acatcgctct tgtacactac	300
actgacggtg tgaaagcata catcatcgct ccaaaaggtc ttgaagtagg tcaacgtatc	360
gtttcaggtc cagaagcaga tatcaaaagtc ggaacgctc ttocacttgc taacatccca	420

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gttggtaactt tgattcaca catcgagttg aaaccagggtc gtgggtgtga attggtacgt 480
gctgctgggtg catctgtctca agtattgggt tctgaaggta aatatgttct tgttcgtctt 540
caatcagggtg aagttcgtat gattcttgga acttgccgtg ctacagttgg tgttgcggga 600
aacgaacaac atggacttgtt aaaccttggt aaagcaggac gttagccgtt gaaaggtatc 660
cgcccaacag ttctgtggttc tgtaatgaac cctaacgatc acccacacgg tgggtggtgaa 720
ggtaaagcac cagttggtcg taaagcacca tctactccat ggggcaaac tgcctctggt 780
cttaaaactc gtaacaagaa agcgaaatct gacaaactta tcgttcgtcg tcgcaacgag 840
aaataa 846

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<210> 439
<211> 366
<212> DNA
<213> Streptococcus pneumoniae

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<400> 439
gcaactagta aatccgccag ctccgttagcg ctccatagga gtgcaagccg ctgtggtaca 60
acatttaag gagaaaatat aaaaatggga cgcagtccta aaaaaggacc tttcgtcgat 120
gagcatttga tgaaaaaagt tgaagctcaa gctaacgacg aaaagaaaaa agttattaaa 180
acttggtcac gtcgttcaac gatcttccca agtttcattg gttacactat tgcagtttat 240
gacggagcta aacacgtacc tgtttacatc caagaagaca tggtaggcca caaacttggt 300
gaatttgcat caactcgtac ttacaaaggt cacgctgcag acgacaagaa aacacgtaga 360
aaataa 366

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<210> 440
<211> 345
<212> DNA
<213> Streptococcus pneumoniae

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<400> 440
atggcagaaa ttacttcagc taaagcaatg gctcgtacag tacgtgtttc acctcgtaaa 60
tcacgtcttg ttcttgataa catccgtggt aaaagcgtag ccgatgcaat cgcaatcttg 120
acattcactc caaacaagc tgcgtgaaac atcttgaaag ttttgaactc agctgtagct 180
aacgtgaaa acaactttgg ttgtgataaa gctaacttgg tagtatctga agcattcgca 240
aacgaaggac caactatgaa acgtttccgt ccacgtgcga aaggttcagc ttcaccaatc 300

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aacaaacgta cagctcacat cactgtagct gttgcagaaa aataa 345

<210> 441

<211> 666

<212> DNA

<213> Streptococcus pneumoniae

<400> 441

ggaggtaaaa tcgtgggtca aaaagtacat ccaattggta tgcgtgtcgg catcatccgt 60
gattgggatg ccaaatggta tgctgaaaaa gaatacgcgg attaccttca tgaagatctt 120
gcaatccgta aattcgttca aaaagaactt gctgacgcag cagtttcaac tattgaaatc 180
gaacgcgcag taaacaaagt taacgtttca cttcacactg ctaaacaccag tatgggtatc 240
ggtaaaggtg gtgctaactg tgatgcactc cgtgcaaaac ttaacaaatt gactggaaaa 300
caagtacaca tcaacatcat ogaaatcaaa caacctgatt tggatgtcga cctgtagggt 360
gaaggaattg ctcgtcaatt ggagcaactg gttgctttcc gtcgtgcaca aaaacaagca 420
atccaaactg caatgcgtgc tggagctaaa ggaatcaaaa ctcaagtatc aggtcgtttg 480
aacggtgcag atatgcccg tgcgtgaagga tactctgaag gaactgttcc gcttcacaca 540
cttcgtgcag atatcgatta cgtctgggaa gaagcagata ctacatacgg taaacttggt 600
gttaaagtat ggatctaccg tggatgaagt cttccagctc gtaaaaacac taaaggaggt 660
aataa 666

<210> 442

<211> 384

<212> DNA

<213> Streptococcus pneumoniae

<400> 442

aaggagaaaa ctgaaatgat tcaaacagaa actcgtttga aagtcgcaga caacagcgg 60
gctcgcgaaa tcttgactat caaagttctt ggtggttcag gacgtaaatt tgcaaacatc 120
ggtgatgtta tcgtggcatc tgtaaaacaa gctactctg gtggtgcgggt taaaaaggt 180
gacgttgta aagcagttat cgttcgtact aaatcagggt ctcgtcgtgc tgatggttca 240
tcatcaaat ttgacgaaaa cgcagcagtt atcatccgtg aagacaaaac tctcgcggga 300
acacgtatct ttggcccagt tgcacgtgaa ttgcgtgaag gtggcttcat gaagatcgtg 360
tcaactgtct cagaagtact ttaa 384

<210> 443
 <211> 327
 <212> DNA
 <213> *Streptococcus pneumoniae*

 <400> 443
 gaaaaatcaa ggagaaacct aatgttttga aaaaaaggcg acaagttcg cgtaatcgtt 60
 ggtaaageta aggaacaga agctgttgc cttactgcc ttccaaaagt aaacaaagtt 120
 atcgttgaag gtgttaacat tgttaagaaa caccaacgtc caactaacga gcttccctcaa 180
 ggtggtatca tcgagaaaga agcagctatc cagctatcaa acgttcaagt ttggacaaa 240
 aatggtgtag ctggtcgtgt tggatacaaa tttgtagacg gtaaaaaagt tcgctacaac 300
 aaaaaatcag gcgaagtgtc tgattaa 327

<210> 444
 <211> 570
 <212> DNA
 <213> *Streptococcus pneumoniae*

 <400> 444
 ttaatcacga aggaaggag aagtataatg gcaaatcgtt taaaagaaaa atatcttaat 60
 gaagtagttc ctgctttgac agaacaattc aactactcat cagtgatggc tgtgcctaaa 120
 gttagataaga ttgttttgaa catgggtgtt ggtgaagctg tatcaaacgc taaagcctt 180
 gaaaaagctg ctgaagaatt ggcacttata tcagggtcaaa aaccacttat cactaaagct 240
 aaaaaatcaa tcgcccgtt cgtcttcgt gaagggtgtg cgatcgggtc aaaagttacc 300
 cttcgtggtg aacgtatgta cgaattcttg gataaattgg tatcagtttc acttccacgt 360
 gtacgtgact tccacggtgt cccaacaaaa tcatttgatg gacgcgggaa ctacacactt 420
 ggtgtgaag aacaattaat ctcccgaa atcaacttcg atgacgttga caaaactcgt 480
 ggtcttgaca tcgttatcgt aacaactgct aacactgacg aagagtcacg tgcattgctt 540
 acaggccttg gaatgcctt tgcaaaaata 570

<210> 445
 <211> 414
 <212> DNA
 <213> *Streptococcus pneumoniae*

 <400> 445
 gagagaaaa acaaaatggt tatgactgac ccaatcgacg acttcctaac tcgtattcgt 60
 atgctaacc aagctaaca cgaagtactt gaagtacgt catcaaacat caaaaaggg 120

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attgctgaaa tccttaaacy cgaagggttt gtaaaaaacg ttgaaatcat tgaagatgac 180
aaacaaggcg tcatacgtgt atttcttaaa tacggaccac atggtgagaa agttatcact 240
aacttgaaac gtgtttctaa accaggactt cgtgtctaca aaaaacgtga agaccttcca 300
aaagtcttta acggacttgg aattgccatc ctttcaactt ctgaagggtt gcttactgat 360
aaagaagcac gccaaaagaa tgttggtggt gaggttatcg cttacgtttg gtaa 414

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<210> 446
<211> 552
<212> DNA
<213> Streptococcus pneumoniae

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<400> 446
caggagaaaa taaacatgtc acgtattggt aataaagtta tcgtgttgcc tgctggtggt 60
gaactcgcta acaatgacaa cgttgtaact gtaaaaggat ctaaaggaga acttactcgt 120
gagttctcaa aagatattga aatccgtgtg gaaggtagtg aaataactct tcacctcca 180
aacgattcaa aagaaatgaa aactatccac ggaactactc gtgccctttt gaacaacatg 240
gttggtggtg tatcagaagg attcaagaaa gaacttgaaa tgcgtggggt tggttaccgt 300
gcacagcttc aaggatctaa acttgttttg gctgttggtt aatctcatcc agacgaagtt 360
gaagctccag aaggaattac ttttgaactt ccaaacccaa caacaatcgt tggtagcgga 420
atttcaaaag aagtagtttg tcaaacagct gottacgtac gtagccttcg ttcaccagaa 480
ccatataaag gtaaagggtat ccgttacggt ggtgaattcg ttgcgcgtaa agaaggtaaa 540
acaggtaaat aa 552

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<210> 447
<211> 426
<212> DNA
<213> Streptococcus pneumoniae

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<400> 447
tcatacaacca ccaacctatt ttccaacttt gtgcatagca cagcatttaa aactaaagag 60
gtgaaaaactg tgatttcaaa accagataaa aacaaactcc gccaaaaacg ccacctgcgc 120
gttcgcggaa aactctcttg aactgctgat cgccacggtt tgaacgtatt cgttctaat 180
acaggcatct acgctcaagt gattgatgac gtagcgggtg taacgctcgc aagtgtctca 240
actcttgata aagaagtctc aaaagggaact aaaactgaac aagccgttgc tgcgggtaaa 300

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ctcgttgacag aacgtgcaaa cgctaaagggt atttcagaag tgggtgttcga ccgcggtgga 360
 tatctatatac acggacgtgt gaaagctttg gctgatgcag ctcgtgaaaa cggattgaaa 420
 ttctaa 426

<210> 448
 <211> 498
 <212> DNA
 <213> *Streptococcus pneumoniae*

<400> 448
 aaaatggcat ttaaagacaa tgcagttgaa ttagaagaac gcgtagtgtc tgtcaaccgt 60
 gttacaaaag ttgttaaagg tggacgtcgt cttcgtttcg cagctcttgt tgttgttggg 120
 gaccacaatg gtcgcgtagg atttggtact ggtaagctc aagaagtcc agaagcaatc 180
 cgtaaacgag tagatgatgc taagaaaaac ttgatcgaag ttcctatggt tggaaacaaca 240
 atccccacacg aagttctttc agaattcggg ggagctaaag tattgttgaa acctgctgta 300
 gaaggtttcg gagttgccgc tgggtggtgca gttcgtgccg ttgtggaatt ggcaggtgtg 360
 gcagatatta catctaaatc acttggttct aacactccaa tcaacattgt tcgtgcaact 420
 gttgaagggt tgaacaatt gaaacgcgct gaagaaattg ctgcccttcg tggattttca 480
 gtttctgatt tggcataa 498

<210> 449
 <211> 468
 <212> DNA
 <213> *Streptococcus pneumoniae*

<400> 449
 cattttacaa aagagagaaa aataaaaatg aaacttcacg aattgaaacc tgcagaagggt 60
 tctcgtaaag tacgtaacgc cgttggtcgt ggtaactcat caggtaacgg taaaacatct 120
 ggtcgtgggt aaaaaggta aaaaagctcgt agcgggtggcg gagttcgcct tggttttgaa 180
 ggtggacaaa ctccattgtt ccgtcgtctt ccaaaaacgtg gattcactaa catcaacgct 240
 aaagaatacg caattgtgaa ccttgaccaa ttgaacgtct ttgaagatgg tgctgaagta 300
 actccagttt ttcttatcga agcaggaatt gttaaagctg aaaagtcagg tattaaaatt 360
 cttggtaacg gtgagttgac taagaaattg actgtgaaag cagctaaatt ctctaaatca 420
 gctgaagaag ctatcactgc taaagggtgt tcagtagaag tcacttaa 468

<210> 450
 <211> 1314
 <212> DNA
 <213> *Streptococcus pneumoniae*

<400> 450
 cctatgtttt ttaaattatt aagagaagct cttaaagtca agcagggttcg atcaaaaatt 60
 ttatttcaaa tttttatcgt ttgggtcttt cgtatcgga ctagcattac agttcctggt 120
 gtgaatgcca atagcttgaa tgccttaagt ggattatcct tcttaaacaat gttgagcttg 180
 gtgtcgggga atgcocctaaa aaacttttcg atttttgccc taggagttag tccctatatc 240
 accgcttcta ttgttgtcca actcttgcaa atggatattt taccacaagt ttgtagagtgg 300
 ggtaacaag gggaagtagg tcgaagaaaa ttgaatcaag ctactcgtta tattgctcta 360
 gttctcgtt ttgtgcaatc tatcgggatt acagctgggt ttaatacctt ggcgtggagct 420
 caattgatta aaactgcttt aactccacaa gttttcttga cgattgggtat catcttaaca 480
 gctgtagta tgattgtcac ttggttgggt gagcaaatca cagataaggg atacggaaac 540
 ggtgtttcca tgattatctt tgcggggatt gttcctcaa ttccagagat gattcagggc 600
 atctatgtgg actactttgt gaacgtccca agtagcogta tcaacttcac tatcattttc 660
 gtaatcattt tgattattac tgtattgttg attatttact ttacaactta tgttcaacaa 720
 gcagaatata aaattccaat ccaatatact aagggtgcac aagggtgctcc atctagctct 780
 taaccttcgt taaaagtaaa cctgctgga gttatcctg ttatctttgc cagttcgatt 840
 actgcagcgc ctgcggctat tcttcagttt ttgagtgcga caggctatga ttgggcttgg 900
 gtaagggtag cacaagagat gttggcaact actctccaa ctggtattgc catgtatgct 960
 ttgttgatta ttctctttac attctcttat acgtttgtac agattaatcc tgaaaaagca 1020
 gcagagagcc taaaaaagag tggtgccatc atccatggag ttcgtcctgg taaaggtaaa 1080
 gaagaatata tgtctaaact tcttcgtcgt cttgcaactg ttggttccct ctctcctggt 1140
 gtgatttcca ttttaccgat tgcagctaaa gatgtatttg gtctttctga tgtgtgtgac 1200
 ttgtgtgaa caagctctct gatcattatc totacaggta tcgaaggaa caagcaattg 1260
 gaaggttacc tattgaaacg taagtatgtt gggttcatgg acagaacaga ataa 1314

<210> 451
 <211> 651
 <212> DNA
 <213> *Streptococcus pneumoniae*

<400> 451
 ggagatcaaa tcatgaatct tttgattatg ggcttacctg gtcagggtaa gggaaactcaa 60
 gcagcaaaaa tcgtagaaca attccatgtt gcacatatct caacagggtga tatgttccgc 120
 gctgcaatgg caaatcaaac tgaatatgggt gttcttgcta agtcatatat tgacaaggggt 180
 gaattggttc ctgacgaagt tacaatatgga atcgtaaaag aacgcctttc acaagatgat 240
 attaaagaaa caggatttctt attggatgggt taccacgta caattgaaca agtccatgcc 300
 ttggacaaaa cattggctga acttggcatt gaactagaag gtgttatcaa tattgaagtg 360
 aacctgaca gccttttggga acgtttgagt gggcgatatca tccaccgcgt aactggagaa 420
 actttocaca aggtctttaa cccaccagtt gactataaag aagaagatta ctaccaacgt 480
 gaagatgata agcctgagac agtaaaacgt cgttttgatg ttaatatgtc tcaaggagaa 540
 ccaatcattg ctactaccg tgccaaagggt ttggttcatt acatcgaagg taatcaagat 600
 atcaatgatg tcttctcaga tattgaaaa gtattgacaa atttgaata a 651

<210> 452
 <211> 378
 <212> DNA
 <213> Streptococcus pneumoniae

<400> 452
 aaaggagaaa acatggctcg tattgtcgga gttgatattc caaatgacaa acgcgtagta 60
 atctcattga cttatgttta tggatcgga cttgcaacat ctaagaaaat ttgggtgct 120
 gctggaatct cagaagatgt tcgtgtacgt gatcttacat cagatcaaga agatgctatc 180
 cgtcgtgaag tggatgcaat caaagtgaa ggtgacctc gtcgtgaagt aaacttgaac 240
 atcaaacgtt tgatggaaaat cggttcatc cgtggatcc gtcacgctg tggacttctt 300
 gtcggtggac aaacactaa aaacaacgcc cgcactcgta aaggtaaagc tgttgcgatt 360
 gctggtaaga aaaaataa 378

<210> 453
 <211> 396
 <212> DNA
 <213> Streptococcus pneumoniae

<400> 453
 gaggtaaaag tcttggttaa accaacacgt aaacgtcgtg tgaaaaagaa tatcgaatct 60
 ggtattgtc atattcacgc tacatttaac aacactattg ttatgattac tgatgtgcatt 120

ggtaatgcaa ttgcttggtc atcagctggt gctcttggtt tcaaagggtc tcgtaaatct 180
 acaccattcg ctgctcaaat ggcttctgaa gctgctgcta aatctgcaca agaacacggt 240
 cttaaatcag ttgaagttac tgtaaaagggt ccagggttctg gtcgtgagtc agctattcgt 300
 gcgcttgctg ccgctggtct tgaagtaaca gcaattcgtg atgtgactcc agtgccacac 360
 aatgggtgctc gtccctccaaa acgtcgccgt gtataa 396

<210> 454
 <211> 402
 <212> DNA
 <213> *Streptococcus pneumoniae*

<400> 454
 ataaaggagg aatacatggc ttaccgtaaa ctaggacgca ctagctcaca acgtaaaagca 60
 atgcttcgctg atttgacaac tgaccttttg atcaacgaat caatcgtgac aactgaagct 120
 cgtgctaag aaatccgtaa aactgttgaa aaaatgatta ctctaggtaa acgtggtgat 180
 ttgcatgcac gtcgtcaagc agctgctttc gtacgtaatg aaatcgcac tgaaaaactat 240
 gatgaagcaa ctgataagta cacttctact acagcacttc aaaaattggt ctcagaaatc 300
 gcacctcggt atgctgaacg taacgggtgga tacactcgta tccttaaaac tgaatcacgt 360
 cgtggtgatg cagcgccaat ggcatcact gaattagtat aa 402

<210> 455
 <211> 414
 <212> DNA
 <213> *Streptococcus pneumoniae*

<400> 455
 atgcctacaa ttaaccaatt ggttcgcaaa ccgcgtaaat caaaagtaga aaaactctaa 60
 tcaccagctt tgaacgttgg ttacaatagt cataaaaaag ttcaacacaa cgtttcttca 120
 ccacaaaaac gtggtgttgc aactcgtgtt ggaacaatga cacttaaaaa acctaactca 180
 gcccttcgta aattcgctcg gtacgttttg agcaacctta tcgaagttag tgctacatc 240
 ccaggatatg gacacaactt gcaagagcac agcgtggtgc ttcttcgctg tggacgtgta 300
 aaagaccttc caggggtacg ttaccatata gtcctggtg cacttgatag tgcagggtgt 360
 aacgatcgta aacaaggccg ttctaaatac ggtactaaac gtccaaaagc ataa 414

<210> 456

<211> 477

<212> DNA

<213> *Streptococcus pneumoniae*

<400> 456

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agagaaatga gtcgtaaaaa tagagctcca aaacgtgacg tattgccaga tccgctatac      60
aatcacaaac tagttactcg tcttatcaac cgcgttatgc ttgatggtaa acgtggtact      120
gctgcttcaa tcgtttacgg tgcttttgag caaatcaaa aagctactgg caacgatgca      180
cttgaagtat ttgaacacgc tatggaaaa atcatgcctg tacttgaagt acgtgcacgt      240
cgtgttggtg gttctaacta ccaagtccca gttaaagttc gtccagaacg tcgtacaaca      300
cttggaactc gttgggttgg aacaatcgct cgtcttcgtg gtgaacacac aatgcaagac      360
cgtcttgcaa aagaaatctt ggatgctgct aacaacactg gtgcagcagt taagaacagt      420
gaagatactc accgtatggc tgaagctaac cgtgcattcg cacacttccg ttggttaa      477

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<210> 457

<211> 930

<212> DNA

<213> *Streptococcus pneumoniae*

<400> 457

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gaaaggactg aaaacatcat gactgaaaaa cttcaattaa ctaaatacga tcgtaaaaaa      60
gtttggtggc gttcaacctt cttacaaggg tcttggaact ttgaacggat gcaaaacttg      120
ggctgggctt atacctcat tccagctatc aaaaaactct atactaaaaa agaagatcaa      180
atcgtctctc ttgagcgtca ccttgagttc ttcaaacactc atccatacgt agctgcteca      240
gtcatggggg ttactcttgc gcttgaagaa gaacgtgcta acggtgtgga aatcgatgac      300
gctgctatcc aagggggttaa aatcggtatg atgggacctc ttgctgggat cggtgaccca      360
gtattctggt ttacagtagc cccaatcctt ggatctctcg gtgcttcaat tgcccttact      420
ggcaatatct tggggccact cctctctctt gttgcatgga acttgattcg tatgcatctc      480
ttgtggtatg ttcaagagat tggatacaag gctggatcag aaatcactaa agatattgct      540
gggtggtatc ttcaagatat cactaaagga gcttctatcc ttgggatgtt cattcttgct      600
gtccttgttc aacgtggggt aaatattaaa ttgctttcgt atgtttctaa agttcaacta      660
gatgaaaagg cttatatcca ttgggataaa ttgccagaag ggtctaaggg tatccaagaa      720
gcattcgcaac agtaggagca aggattgtct caaactcctg aaaaagtac tactttccaa      780
caaaacttgg atatgttgat tcttgatta tcaggactac tcttactttt actttgcagt      840

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tacttactta agaaaaaagt atctccaatc actattatcc ttgcctctct cgcagtgagg 900
 attgtggcac atgttcttca catcatgtaa 930

<210> 458

<211> 831

<212> DNA

<213> Streptococcus pneumoniae

<400> 458

gccattatct atgaaaggat tttaaacatg tctattatct ctatgggttt agtagtcgtt 60
 gtagccttct ttgcagggtct tgaaggcatc ctgcaccagt tccaatttca ccaaccactt 120
 gtagcctgta cccttattgg gcttctaaca ggctcacttg aagcagggat tatcctcggt 180
 ggatcgcttc aaatgattgc ccttggttgg tcaaatatcg gtgctgctat cgctcctgat 240
 gctgcacttg cttctgtcgc tgcggcatt atcatgggtc ttggtgggtga ctttaaccaag 300
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 acaatgattg ttctgacaat ttcagtttgt ttggttcata ctgcagatgc tgcgcgtaaa 420
 aaagggtgact tcggcgctgt ggagcgtgcg catttcacgc cgctactttt ccaaggactt 480
 cgtatcgctc ttctgcgcgc tcttctcctt atgggtaccaa ctgaaactgt acaaagtatc 540
 cttagtacca tgccagactg gctcaaagat ggtagggcta tcgggtgggt tatgggtcgtt 600
 gccgttggtt acgccatggt tatcaacatg atggcaactc gtgaagtatg gccattcttc 660
 gctcttggtt tcgttctcgc tgcgtgtgta gatattactc taatcggatt cgggtgctatc 720
 ggctgtgcta tcgctcttat ctaccttcac ctttctaaaa ctgggtgaaa tgggtggcgga 780
 ggagccgcaa cttctaacga ccaatcggc gatctcctag aagactacta a 831

<210> 459

<211> 1014

<212> DNA

<213> Streptococcus pneumoniae

<400> 459

aacaaaaag gaggaatgac aataatgagt atcggaatca ttattgcgag ccacggcgaa 60
 tttctgcggc gtattcatca gtcaggatct atgatctttg gtgaacaaga aaaggttcaa 120
 gttgtgacct ttatgcaaaa tgaaggctct gatgatctat acgctaagtt taataacgct 180
 gttgctgcac ttgacgcaga agatgaggtt ctagttttgg ctgacctttg gagtgggttc 240

ccatttaacc aagctagtcg cgtgatggga gaaaatccctg agcgtaagtt tgcctatc 300
acaggactta acttacogat gttgattcaa gctacacag agcgccctcat ggacgtgtgt 360
gcagggtgtg aaaaagtcgc tgctaataatc attaagaag ccaagatgg catcaaagct 420
cttcagaag agctaaatcc agtcgaagaa gttgcaagcg ctgcagctgc tccagttgcc 480
caaatgtcta tccagaaggg aactgttacc ggagacggta aattgaaaat caatcttgcc 540
cgtcttgaca cagctctact tcacggtcag gttgcaactg cttggactcc agattcaaaa 600
gcaaatcgta tcactgttgc ttcagataac gtggctaaag acgaccttcg taaagaattg 660
attaacaag cagctccagg taatgtcaag gctaacgtgg ttccaattca aaaactgatt 720
gagatttcaa aagaccacg ttttggagaa acacatgcc ttatcttgtt tgaacacct 780
caagatgcc tctgtgccat cgaaggcggc gtgccaatca agactcttaa tgttggttct 840
atggctcact caacaggtaa aacattggtc aataccgttt tgtctatgga caaagaagac 900
gttgctacat ttgaaaaat gcgtgacttg ggtgttgaa ttgatgtccg taaagtacca 960
aatgattcta aaaaagattt gtttgacttg attaacaag ccaatgtcaa ataa 1014

<210> 460

<211> 1128

<212> DNA

<213> *Streptococcus pneumoniae*

<400> 460

tatacttcac atataaatac aagagagctc gtcacactcg actctgttga caaaaaatca 60
atccagtcgc gttgtccctg ttctcggtca atatcaagga ggatagctat gaaagctgtt 120
gttgtaaata cagaagcac tgggtgtgct attgaagaaa aagtactccg tccactgaa 180
actggggaag cactttaga agttgaatac tgtggcgttt gccacacoga cctccacgtt 240
gctcatggtg acttttggtca agtcccagga cgtgttcttg ggcacgaagg tatcggatc 300
gttaaagaga ttgtccaga tgtgaaagc cttaaagtcg gtgaccgcgt cagcgttgct 360
tgggtctttg aagtagtggt cacttgcgaa tactgtacaa ctggctcgga aaccccttgc 420
cgtacagtga aaaaatgctg ctactcagta gacggtggtg tggctgaaca atgtatcgta 480
actgtgact atgtgtgcaa agttctgtac ggaactgtac cagcccaagc ttctctatc 540
acatgtgctg gagtacaac ctataaagct atcaaagaag caaaagtga accaggccaa 600
tgggtgtgtc tttacgggtc tgggtgactt ggtaacctcg ctgttcaata cgctaaaaaa 660

gtattcaatg ctcatgttat cgcagtcgat atcaacaatg acaaacttgc ccttgcaaaa 720
 gaagtaggcg ctgacattgt gattaacggc ctcgaagttg aagatgtagc tggactcatt 780
 aaagaaaaaa ctgatggagg agctcattca gctgtcgtaa ctgctgtgtc taaagttgcc 840
 ttcaaccagg ctgttgactc cattcgtgct ggtggtcgag tcgtcgtgtg tggctcttct 900
 tctgaaatga tggaaactcag catcggttaa acagtcctcg atggaatcca agtcacggtg 960
 tctctgtgag gaactcgtaa agacttagaa gaagccttc aatttgggtg agaaggtctg 1020
 gtagtcccag ttgttcaaaa acgtccagta gaagatgctg ttgccatttt cgacgaaatg 1080
 gaaaaaggcc aaatccaagg acgtatggta ctgacttca cccactaa 1128

<210> 461
 <211> 477
 <212> DNA
 <213> *Streptococcus pneumoniae*

<400> 461
 acaaaaatcg aactatatat aggagaaatc atgaacaaaa caacatttat ggctaaacca 60
 ggccaagtgt aacgtaaatg gtacgtagtt gacgcaactg atgtaccact tggacgtctt 120
 tctgcagtag ttgctagcgt acttcgcgga aaaaacaaac caacatttac accacacact 180
 gatacagggtg acttcgtgat tgttatcaat gctgaaaaag ttaaatgac tggtaaaaaa 240
 gcaactgata aaatctacta cactcactca aaccaccag gtggattgaa acaaatctct 300
 gcagggtgaac ttcgttctaa aaatgcagta cgtttgattg agaaatcagt taaaggatg 360
 cttccacaca atacacttgg acgcgctcaa ggtatgaagt tgaagattt tgttggagct 420
 gagcacactc acgctgcaca acaaccagaa gttcttgaca ttccaggact tatctaa 477

<210> 462
 <211> 396
 <212> DNA
 <213> *Streptococcus pneumoniae*

<400> 462
 agtatgtcac aagcacaata tgcagggtact ggacgtcgta aaaacgctgt tgcacgcgtt 60
 cgccttgttc caggaaactgg taaaatcact gttaaacaaa aagatgttga agagtacatc 120
 ccacacgtg accttcgtct tgtcatcaac caaccattcg cagttacttc aactgtagggt 180
 tcatacgacg ttttcgttaa cgttataggt ggtggatacg ctggtcaatc aggagctatc 240
 cgtcacggta tcgtcgtgac ccttcttcaa gtacaccag acttcgcgga ttcattgaaa 300

cgcgaggac ttcttacacg tgactcacgt aaagttgaac gtaagaaacc aggtcttaag 360
 aaagtcgta aagcatcaca atttagtaaa cgtaa 396

<210> 463
 <211> 834
 <212> DNA
 <213> Streptococcus pneumoniae

<400> 463
 aggaggagc acacgatgac tggatctaac aaattaacaa aacgtgatta tottaaaacg 60
 tctttgcggg cattcttttg tcaaaatgga tttaactata gtaactatca aggggtggga 120
 tatgccaatg tgatgtatcc tgccttgaaa aaacactatg gagaggatca ggaagggttc 180
 taccaagcct tgggaagaaa ctgtgaattc tataatacca acccacactt cctgcctttt 240
 attaccagct tgcactcttg aatgttgaa aatggcgtc cgcaaaaga aacacgtagc 300
 atcaagatgg ccttgatggg accattggca ggtattggg attctctttc tcaattctgt 360
 ttagctccat tgttctcaac catgcgagct tcgtttgtct aagaaggctt ggttgcgggt 420
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 gttaagatta tgggtccgat tacctttgct gcaggggaag ttaaagcaga cgctaataca 660
 agtatcgtaa gtattcaggg aatgcttgat aaggttgtct cagctcttct accagcccta 720
 tttaacttt tagtttacta ctgcatcaaa gaaaagaat ggacaacata taaactcggt 780
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<210> 464
 <211> 1533
 <212> DNA
 <213> Streptococcus pneumoniae

<400> 464
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 caatttgacg gcaagctcta cttagagttt ggcggtaaaa tgtagaaga tttccacgct 180
 gctcgtgtcc ttctcggtta tgaacctgac aacaaatca agctcttgca agaattgaaa 240

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tactaccaat acgaccgctt atatgtaag taa 1533

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<210> 465

<211> 2166

<212> DNA

<213> Streptococcus pneumoniae

<400> 465

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attccacag atttgggttaa ggcaatogtt tctatcgaag accatcgctt cttcgaccac	300
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cctgctccac aacaaccccc atcaactgaa agttcaagct catcatcaga tagttcaact	2040
tcacagteta gctcaaccac tccaagcaca aataatagta cgactaccaa tcttaacaat	2100
aatacgcaac aatcaaatc aaccctgat caacaaaatc agaatcctca accgacgcaa	2160
ccataa	2166

<210> 466
 <211> 351
 <212> DNA
 <213> Streptococcus pneumoniae

<400> 466	
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caagagtttg gacgtgaggt cgtgggtat aataaagtag aagttgacga gtttttagac	120
gatgtcatca aggactatga aaactatgct gccttggtca agtcacttcg tcaggaaatt	180
gcggtattga aggaagaatt aactcgtaaa cgaacacctt caccagttca agcagaaccc	240
cttgaagcgg caattacaag ttctatgacg aattttgata ttttgaaacg cctgaataga	300
ttgaaaaaag aagtttttgg taaacaaatt ttagataact cagattttta a	351

<210> 467
 <211> 1407
 <212> DNA
 <213> Streptococcus pneumoniae

<400> 467	
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caatttgatt ttgatgaagc aaaagagcta acagttgggtc aagctattcg taaaaatgaa	120
gaagtggaat caggagtctt gcctgaggat tccattttgg acaagtatgt taagcaacac	180
agagatgaaa ttgaggcgga taagtttgcg actogtcaat acaaaaaaga ggagttcgtt	240
gaaactcaga gtctggatga tttaattcaa gagatgcgtg aggcgtgtaga gaagtcagaa	300
gcttcttcgg aggaagtctc atcttctgaa gacatcttac tacccttgcc tctggacgat	360
gaggagcaag gcttgatgcc tctattgcta gatgatgaaa atccaacaga aatgactgaa	420

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gaagtggaag aggagcaaaa cctttctcgt ctggatcaag aggactcaga aaagaaaagt 480
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tattatgtct accgtcaagt ggctcgttcg actaaggaaa ttgaaacttc tcaatcaact 600
acagccaatc aatcggatgt ggatgatttt aatacacttt atgacgcctt ttacacagat 660
agcaataaaa cggccttgaa aaatagccag tttgataaac tgagtcaact caagacttta 720
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tatatccttg aactgttcaa tatcgttaac ggcaatggtt attacaacct ctacaagcca 1320
gatggaacct atctctttac ccttaactgt aagacaggct actttgtcgg aaatggcgct 1380
ggtcatgcgg atgacttaga ttactaa 1407

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<210> 468
<211> 1500
<212> DNA
<213> Streptococcus pneumoniae

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<400> 468
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cgtaaccttg cccttaatat tgaatctcgt ggttacacag ttgctatctc caaccgtagt 180
aaagaaaaaa cggaagatgt gattgcttgc catcctgaaa agaactttgt accaagctat 240
gacgttgaaa gttttgtaaa ctcaatcgaa aaacctcgtc gtatcatgct gatgggttcaa 300
gctggacctg gtacagatgc tactatccaa gcccttcttc cacaccttga caagggtgat 360
atcttgattg acggaggaaa tactttctac aaagatacca tccgtcgtaa tgaagaattg 420

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gcaaactctg gtatcaactt tatcggtact ggggtttctg gtggtgaaaa aggtgccctt 480
gaagtgccct ctatcatgcc tgggggacaa aaagaagcct acgaattgggt tgcggatggt 540
cttgaagaaa tctcagctaa agcaccagaa gatggcaaac catgtgtgac ttacatcggt 600
cctgatggag ctggtcacta tgtgaaaatg gttcacaatg gtattgagta cgggtgatatg 660
caattgatcg cagaaagcta tgacttgatg caacacttgc taggcctttc tgcagaagat 720
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cgttcagctg accttcacgc taacttgatc caagcacaac gtgactactt tgggtgctcac 1440
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<210> 469

<211> 438

<212> DNA

<213> Streptococcus pneumoniae

<400> 469

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acgattgatg tcatcggtaa ggctccagac gtgactccaa gtcaagtgtc aaaagagttg 180
atggtaactc ttggaactgt tacgacaagt ttgaacaatt tagagcgtaa gggctacatt 240
gagcgagttc ggtcagaaca ggatcgtcgt gtgggtgcatc tgcatttgac aaagaagggt 300
cgcttgatgc atagactgca taaacgcttc cacaaggcca tggtagaaaa aattattgat 360

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ggcattgagcg aggaagaat tgctgtcatg ggtaaagggt tgactaatct ttaccaattt 420
 ttggaggatt tgaataaa 438

 <210> 470
 <211> 237
 <212> DNA
 <213> Streptococcus pneumoniae

 <400> 470
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 aaagacgcatt cagaagtaac acttgaatca acttttgatg atttggcagc agattcattg 120
 gacttggtcc aagtaatctc agaaatcgaa gatgcttttg atatccaaat cgaagcagaa 180
 aatgacttga aaacagttgg tgacttggtt gcttacgttg aagagcaagc aaaataa 237

 <210> 471
 <211> 2343
 <212> DNA
 <213> Streptococcus pneumoniae

 <400> 471
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 aaagcttggg aaggcttcaa aggcgttagat tggaaagaaa aagcaagtgt atcaccagttt 120
 gtacaagcta actacacacc ttatgatgga gacgaaagct tccttgacag accaacagag 180
 cgttcacttc acatcaagaa aattgtagaa gaaactaaag cactactcga agaaactcgt 240
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 gttaacattg ctctcatggc tgtctgcgtg gtgattaacg gtgctgctac atctctaggt 840
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gaatcagaaa tccaagaatt cyttgatgat ttcgttatga aacttcgtac agttaaat 960
gctcgtacaa aagcttatga ccaattgtac tcagggtgacc caacctttat cacaacttct 1020
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taa 2343

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<210> 472
<211> 1620
<212> DNA

<213> *Streptococcus pneumoniae*

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 gatgtcatta tcacagaggt tgggtggaaca gtaggagata tcgagtcctt gccattccta 480
 gaggctcttc gtcagatgaa ggcagatgtg ggtgcggata atgtcatgta tatccataca 540
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 gtcaagaat tgcgtggctt gggaatcaa ccaatatgt tggttattcg tacagaagag 660
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 gttatcgaat cgttgatgt tgaacacctt taccaaatc cactgaactt gcaggcacaa 780
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 gttggttaagt atgtggagtt gcaagatgcc tatatctcag tggtcgaagc cttgaaacac 960
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 attcctgaaa ataaattctt ttagcttgt cagtatcacc ctgaactgtc aagccgtcca 1560

aaccgaccag aagaactcta cactgccttt gttactgcag cagttgagaa cagcaattag 1620

<210> 473

<211> 1140

<212> DNA

<213> Streptococcus pneumoniae

<400> 473

attatgaaca atactgaatt ttatgatcgt ctgggggtat ccaaaaaacgc ttccggcagac 60

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cctggtgctg aggacaagta caaggaagtt caagaagcct atgagacttt gagtgcagac 180

caaaaacgtg ctgcctatga ccagtatggt gctgcaggcg ccaatggtgg ttttggtgga 240

gctggtggtt tcggcggttt caatggggca ggtggcttcg gtggttttga ggatattttc 300

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actcctcttg gtatgatgcy tcgccaagta acctgtgatg tctgtcacgg tcgaggaaaa 600

gaaatcaaat atccatgtac aacctgtcat ggaacaggto atgagaaaca agctcatagc 660

gtacatgtga aaatccctgc tgggtgtgaa acaggtoaac aaattcgctc cgctggtcaa 720

ggtgaagcag gctttaacgg tggaccttat ggtgacttgc atgtagtagt ttctgtggaa 780

gctagcgaca agtttgaacg tgaaggaaac actatcttct acaatctcaa cctcaacttt 840

gtccaagcgg ctcttggtga tacagtagat attccaactg ttcacggtga tgttgaattg 900

gttattccag agggaaactca gactggtaag aagttccgcc tacgtagtaa gggggcacgc 960

agccttcgtg cgggtgaggt tggtagacca tacgttactg ttaatgtcgt¹ aacaccgaca 1020

ggcttgaacg accgccaaaa agtagccttg aaagaattcg cggctgctgg tgacttgaaa 1080

gtaaatccaa agaaaaaagg cttctttgac catattaaag atgcctttga tggagaataa 1140

<210> 474

<211> 297

<212> DNA

<213> Streptococcus pneumoniae

<400> 474

ggagttgcta tgaattatc caacctactg ctatttgacg gagctgcagc cggaagttat 60

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ctggttacaa aaaatcgcca aaccatcaca gatgaagtct tgaataccac tgaccgcgtt      120
caagctatca aggacgatgt ggatattatc caaaacagcc tgcaaatcat taaccagcaa      180
aaagaactta tcaaggaata ccaagaagac ttgacttaca agtttaaggt cttggaaaag      240
gatatccaga ctgactagc tgtgataaaa gaaatgcagg gaactgaaga taagtaa          297

<210> 475
<211> 1158
<212> DNA
<213> Streptococcus pneumoniae

<400> 475
ggattcaaaa gtgaagaaaa catgagtaaa gaaatgctag aggccttcgc cattttggaa      60
gaagacaagg gaatcaaaaa agaagatata atcgacgcag tagtagagtc gcttcgttcc      120
gcttatcgca gacgctatgg tcagtcagac agcgtagcta ttgacttcaa cgaaaaaaca      180
ggtagcttta cagtttatac tgtccgtgaa gttgttgatg aagtatttga tagccgtttg      240
gaaatcagct tgaagatgc tcttgccatt aattcagctt atgaacttgg agacaaaatc      300
aagtttgaag aagcaccagc tgagtttggt cgtgtagcag cccaatctgc caaacaacc      360
atcatggaaa aaatgcgcaa gcaaacacgt gccatcactt acaacttta caaagaacat      420
gagcaagaaa tcatgtctgg tacagtagaa cgctttgaca accgctttat ctatgtcaac      480
cttggtagca tcgaagccca attgtcaaaa caagacccaa ttccctggaga agtttttgc      540
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gaagtttatg atggaactgt tgaatatcat agcgtggctc gtgaagcagg tgaccgtacg      720
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ggtggtgcta atatcaagaa gattactagc aaattccacc cagctcgtta cgatgctaaa      840
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aacgacagca aacgtgcctt ggtggttggt ccagataaca agctttctct tgccattgggt      1020
cgtcgtggac aaaacgtgag cttggcggct cacttgactg gttaccgtat cgatatcaag      1080
tctgctagcg aatttgaagc catggaagac gctgcttcag tagagttgga agtagaaaaac      1140
gatactgtag aagaataa                                1158

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<210> 476
 <211> 2910
 <212> DNA
 <213> *Streptococcus pneumoniae*

<400> 476
 ggtctcttat ggaatagaag aggaggacat gatttgtcta agaaaagatt gtacgaaatc 60
 gcaaaagaac ttggaaaaga aagtaaagaa gttgtagcgc gtgcaaaaga gttgggcttg 120
 gatgtgaaaa gccactcatc aagtgtggaa gaagctgtcg ctgcaaaaat tgcgtccagc 180
 tttaagcctg cagctgtctc gaaagtagaa gcaaaacctg cagcccaaaa agtaagtgtca 240
 gaaaagaaag ccgaaaaatc tgagccagct aaaccagctg tagctaagga agaggcaaaa 300
 cctgcagccc caaaagcaag tgcagaaaag aaagccgaaa agtctgaacc agtaaaacca 360
 gctgtagcca aggaagaggc aaaaccagct gagccagtc aatccgaaaa agaaaagta 420
 ggggtataac cgcaaatgtg taatttcaag gctgagcgtg aagcacgtgc taaagagcag 480
 gcagagcgac gcaagcaaaa taagggcaat aaccgtgacc aacaacaaa cggaaaccgt 540
 cagaaaaacg acggcgttaa tgggtgaaaa caaggtcaaa gcaaccgca caatcgtcgc 600
 ttaaatgacc aagctaagaa gcagcaaggt cagcaaaaac gtagaatat gcgccgtcag 660
 caagagata aacgttcaaa tcaagcggct ccacgtattg actttaaac ccgtgcagca 720
 gccctaaaag cagagcaaaa tgcagagtac gctcgttcaa gtgaggaaac cttcaagcag 780
 tatcaggctg ctaaaagaac cttggctcaa gctaacaac gcaaggaaac agaggaaatc 840
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 aaaaatcgtg acgattatga tcatgaagaa gatggtccta gaaaacaaca aaagaatcga 1020
 agtagtcaaa atcaagttag aaatoaaaag aatagtaact ggaataacaa caaaagaac 1080
 aaaaaggca atacaagaa caaccgtaat cagactccaa aacctgtac ggagcgtaaa 1140
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 gccacacaaa accaatcett ggaatgggaa acaattgaac tctcatggt ggattacggt 1320
 atcgaagcca acaaaaaggt tgaagtggat aatgctgaca tcgaacgttt ctttgcgaa 1380
 gatggttatc tcaatgaaga tgaattgggt gagcgtccac cagttgttac tatcatggga 1440

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cacgttgacc acggtaaaac aacccttttg gatactcttc gtaactcacg tgttgcgaca 1500
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aagattacct tccttgatac accaggacac gcggccctta catcaatgcg tgcgcgtggt 1620
gctcttgta ccgatattac gatcttggtc gtagcggcag atgacggggt tatgcctcag 1680
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gatggtgaac tcgcaagctt gaaacactat aaagacgacg tgaaagaagt gacaaacggg 2820
cgtgaagggt gattgatgat cgacggctac aatgatatta agatggatga tgtgattgag 2880
gcgtatgtca tggaagaaat caagagataa 2910

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<210> 477
<211> 912
<212> DNA

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<213> *Streptococcus pneumoniae*

<400> 477
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 caagcagccc gtgacaacgg ttatgcagtt ggtggattta acacaacaa ccttgagtgg 120
 actcaagcta tcttgcgccg agcagaagct aaaaaagctc cagttttgat ccaacttca 180
 atgggtgctg ctaaatatcat ggggtgttac aaagtgtgct gcaacttgat cgctaacctt 240
 gttgaatcaa tgggtatcac tgtaccagta gctatccacc ttgaccacgg tcaactacgaa 300
 gatgcacttg agtgtatcga agttggttat acttcaatca tgtttgacgg ttcacacctt 360
 ccagttgaag aaaaccttaa attggctaaa gaagttgttg aaaaagcaca cgctaagggt 420
 atctcagtag aagctgaagt tggactatc ggtggtgaag aagatggaat catcggtaaa 480
 ggtgaattgg ctccaatcga agacgctaaa gcaatggtg aaactggtat cgactctctg 540
 gcagctggta tcggtaacat ccacggtcct taccagtaa actggaagg tcttgacctt 600
 gaccacttgc aaaaattgac agaagctctt ccaggattcc caatcgtatt gcacggtgga 660
 tcaggtattc ctgatgagca aatccaagca gctatcaaac ttggtgtgtg caaagttaac 720
 gtaacacag aatgccaat cgcattcgct aacgcaactc gtaaatgtg tcgtgactac 780
 gaagcaaacg aagcagaata cgacaagaaa aaactcttcg acccagtaa attctggct 840
 gacggtgtaa aagctatcca agcatcggtt gaagaacgta tcgacgtatt cggttcagaa 900
 ggtaagcat aa 912

<210> 478

<211> 468

<212> DNA

<213> *Streptococcus pneumoniae*

<400> 478
 aaatctctaa ttaccgcca aaccacaag gaggatttaa aaatggctaa aaagtcgaa 60
 aaactgttaa aattgcaaat cctgctggt aaagctacac cagctccacc ggttgacct 120
 gctcttggtc aagctggtat caacatcatg ggattcaca aagagttcaa cgcctgtaca 180
 gctgaccaag ctggtatgat cattccagtt gttatctcag ttacgaaga taaatcattt 240
 acttctgta cgaaaacacc accagctgct gttcttttga aaaaagctgc aggtgttgaa 300
 aaagatcag gtacacctaa taaaactaaa gttgctacag ttactcgtgc acaagtacaa 360
 gaaattgcag aaactaagat gccagatttg aacgcagcaa acgtagagtc tgcaatgcgt 420

atgatcgaag gtactgctcg ttctatggga ttcactgttg ttgactaa 468

<210> 479

<211> 693

<212> DNA

<213> Streptococcus pneumoniae

<400> 479

aaaatggcta aaaaagcaa acaacttcgt gctgctcttg agaaaatoga cagcacaaaa 60

gcatacagtg tagaagaagc tgtagcactt gcaaaaagaaa ctaactttgc aaaatttgat 120

gcaactgtag aagttgctta caacttgaac atcgacgta aaaaagctga ccaacaaatc 180

cgtggagcaa tgggtattgcc aaacgggtact ggtaaaactt cacgtgttct tgttttcgca 240

cgtggtgcaa aagctgaaga agcaaaagct gctggtgcag actttgttgg tgaagatgac 300

cttggtgcta aaatcaacga cgggtggttg gacttcgacg tagttatcgc tacacctgat 360

atgatggctc ttgttggacg tcttggacgt gtccttggac cagctaactt gatgccaaac 420

cctaaaactg gtactgtaac aatggatgtt gctaaagcag ttgaagagtc taaagtggtt 480

aaaatcactt accgtgtcga ccgtgcaggt aacgttcaag caatcatcgg taaagtatca 540

tttgaagctg aaaaattggt tgaaaacttt aaagctttca acgaaacaat ccaaaaagca 600

aaaccagcta cagctaaagg aacttacgta acaaaactga ctatcacaa tactcaaggt 660

gttggtatca aagttgacgt aaactcactt taa 693

<210> 480

<211> 1866

<212> DNA

<213> Streptococcus pneumoniae

<400> 480

attttgaata ttatagagga aatcatgaca aaattaagag aagatatccg taacattgcg 60

attatcgccc acgttgacca cggtaaaaca accctggttg acgaattatt gaaacaatca 120

gaaacgcttg atgcacggac tgaattggca gagcgtgcta tggactcaaa cgatatcgaa 180

aaagagcgtg gaattaccat ccttgctaaa aatactgcg ttgcttacaa cggaactcgt 240

atcaacatta tggacacacc aggacacgcg gactttggtg gagaagtga gcgtatcatg 300

aaaatggttg acggtgttgt cttggtcgta gatgcctatg aaggaaccat gccacaaact 360

cgttcgtat tgaaaaaagc cttggaacaa gacottgtcc caatcgtggt tgttaacaaa 420

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atcgataagc catcagctcg tccagcagaa gtagtggatg aagtcttggg acttttcac 480
gagcttgggtg cagatgacga ccagcttgat ttcccagtg tttatgcttc agcgatcaac 540
ggaaacttctt cattgtcaga tgatccagct gaccaagaag cgactatggc accaatcttt 600
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caagtgtcac ttttgacta caatgacttc gttggacgta tcggtatcgg tcgtgtcttc 720
cgtggtacag ttaaggttgg ggaccaagtt accctttcta aacttgacgg tacaactaaa 780
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gaccaactg attcaccaga taaatggact gtttcaggac gtggagaatt gcacttgta 1140
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cttgtttcta tcgatgctgg taaggctaca acttactcaa tcatgtctat cgaagaacgt 1560
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tctcttgagt tcttgaaaga cgatgagtac atggaagtaa cgcttgagtc tatccgtttg 1800
cgtaaacaaa tccttaacaa agcagagcgt gagaagacta acaagaagaa aaaatcagct 1860
gaataa

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<210> 481

<211> 1200

<212> DNA

<213> *Streptococcus pneumoniae*

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<400> 481
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ttggagccaa tggcgcaaca agttattcct aaagcagcat ttggctatat cgctagtgagg      180
gcggaagata ctttcacttt aagagagaat atccgtgcct ttaaccacaa gctcatcggt      240
cctcacacac tttgcaatgt agaaaatcca agtacagaga ttgaatttgc aggtgaaaaa      300
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gaagtggcga ctgcgcgttg tgtgcatgag tttggttctc ttataacaac cagttcttac      420
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ttttacttta gtaaggatga cggatcaaac cgccacatca tggaccgtgt gaaggctgaa      540
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aagacagtta tgcaattgtc tggaaactcag accattgaag atgtcaaaac cttcaagctc      1140
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```

<210> 482
<211> 1011
<212> DNA
<213> Streptococcus pneumoniae

```

```

<400> 482
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gccaaagtac gtgatgagtt tggctacgac atcccaagtg aaaaatcgg agaattattg      180

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gccatctcag cccatccaaa tggagtctct aaagttgcc aatggctgta cgagggaaca 240
gatcttgcta ctttgtatgc ggaacacggt gaattatttg gcaatcgctc agaacctgta 300
tttccacttt tgaccaagat cctcgatgcc aacgactggc tcagtgtcca agttcaccca 360
gacgatgctt atggactcga gcatgaaggg gaactcggaa aacagaaatg ctggtacatt 420
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<210> 483

<211> 864

<212> DNA

<213> Streptococcus pneumoniae

<400> 483

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caactgccaa caccagctat taacacaacc tttgtaagtg tcccttaac cattcaagta 120
gctgatcagg tctttgttga tgacgccagt ctcgatatgt accaaatgat ggaaccatg 180
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gaaactggga ctctogaatt gctacaaaaa gcaaggggat caaagaaatc agttcaagct 660
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 aataacgaaa aatgttgtca acagctctca gagcgaatcc gtgaaacctt cccacaagcg 780
 gatatataaa ttctaccaac ctctgggtctc tgcagtttct atgcagaaga gggcggtttg 840
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<210> 484
 <211> 681
 <212> DNA
 <213> Streptococcus pneumoniae

<400> 484
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 acttctacaa aagcttttctg tgagctagta gatgagattg caatgttgat ggggtatgaa 180
 gtacttcgtg atcttccact agaagatgtg gaaatcgaaa caccaattac aaaaacagtt 240
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 cgtgatgaag aaacacttca accagtgaa tacttggtga aattgcttga ggacattgac 420
 caacgtcaaa tttttgtagt agaccctaat ttggcaacag gtggctcagc aatcttggtc 480
 gttgattctc ttaaaaaacg tggcgcatca aatatcaaat ttgtctgcct tgtatctgct 540
 ccagaggtg taaaagccct tcaagaagct catccagatg tagaaatctt tacagcagcc 600
 ttggatgaac gtttgaacga acacggttat atcgttcag gtcttgaga tgcaggagac 660
 cgcttggtcg gtacaaaata a 681

<210> 485
 <211> 951
 <212> DNA
 <213> Streptococcus pneumoniae

<400> 485
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 tctgaacag ctgaggtgc ttcagttaca actaaccttg taaccaatc taaagatatca 120
 gcagtcgtag gacctgcgac atctgggtgc actgcagctg cggtagcgaa cgctacaaaa 180
 gcagggtgtc cattgatctc accaagtgcg actcaagatg gattgactaa aggtcaagat 240

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aaaggggattg caaaatcttt ccgcgagtc tacaaggggtg aaatcggtgc agatgaaact 420
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aaaggtgcta aaaattcagg tgaatcaag aataacctg ctaaaacaaa agattttgaa 840
gggtgaactg gtcaacaacg cttcgatgca gaccacaaca cagtcaaaac tggttacatg 900
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<210> 486

<211> 1218

<212> DNA

<213> *Streptococcus pneumoniae*

<400> 486

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gataccattg cagagattgg ttataccaat acagaatatg gattttctgc tgagacggtg 300
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aaagaagcag taaaataa 1218

```

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<210> 487
<211> 291
<212> DNA
<213> Streptococcus pneumoniae

```

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<400> 487
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gaaacagttg gaacttacaa ccacttgtt gctgaaaacc aagtaacttt gaaagaagac 180
cgcgttcttg catggttggc taatggagct caaccttcag acacagtacg taacatcctt 240
tcaaaagaag gcgtattgaa aaattccac gattctaaat ttcaaaaata a 291

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<210> 488
<211> 414
<212> DNA
<213> Streptococcus pneumoniae

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<400> 488
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gataagattt accaagccat ctggaaggca gcccaaaactg tttatgtttt gacagatgat 180
ttcgctcaaa atcttctcta agtcactaag aaggtagttt tggatttaca agaagccaag 240
gtggaacgtg cgactatcag tatgattcaa tctatggttg aacatcgttt attgggcgca 300
ggttacatta ccattgcaga acactacatt tctatcggtt tacaacgtga cttggaaaga 360

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agtgggttatg gagatcatat cgcagttcat ttacattttg aacaaattcg ctaa 414

 <210> 489
 <211> 2268
 <212> DNA
 <213> Streptococcus pneumoniae

 <400> 489
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 aatctcaatg gaaaacaaaa acaaatgac ctctgtcaaa actgctataa gattatcaaa 120
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gaatacatgg aaaaacatag tgtggctaag ttggtcggcg ctccctccagg ttatgttggc 1620

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gtcgagctca tgctagcaga tgtaacaag cgccctctcta gcaacaacat tcgttttggat 2040

gtaactgata aggtcaagga aaagttggtt gacctaggtt atgatccaaa aatgggagca 2100

cgccacttc ctgaggactat tcaagactat attgaggaca caatcactga ctactacctt 2160

gaaaatccaa gcgaaaaaga tctcaaaagca gttatgacta gcaagggaaa cattcagatt 2220

aaatctgcc aaaaagctga agttaaaagt tctgaaaaag aaaaataa 2268

<210> 490

<211> 1203

<212> DNA

<213> Streptococcus pneumoniae

<400> 490

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gttgaagggtg tcttgactct tcgcgaattg acaaacgacg gtgatgcaga tatcaatgac 180

tttggttaag taggagaagt attggtgtt ctgtacttc gtcaagtagt tggttaagat 240

actgatacag ttacatacct tgtatctaaa aaacgccttg aagctcgcga agcatgggac 300

aaacttgtt gtcgcgaaga agaagttgtt actgttaaa gaaacgcgtg cggttaagggt 360

ggactttcag tagaatttga aggtgttctg ggatttatcc cagcttcaat gttggatact 420

cgtttcgtac gtaacgtgta gcgttttcta ggtcaagaat ttgatactaa aatcaagaa 480

gttaacgcta aagaaaaccg ctctcatcct tcacgtcgtg aagttgttga agcagctact 540

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taa 1203

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<210> 491
<211> 1443
<212> DNA
<213> Streptococcus pneumoniae

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<400> 491
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ggaaaaggac tcaacgaagg tgttattcgt gaattatctg ctgctaaggg tgagcctgag 180
tggatgttgg agttccggtt gaagctctat gaaaccttca aaaaatgcc catgcaaaact 240
tggggagcag acttgctaga gattgacttt gatgacttaa tctactacca aaacacatct 300
gacaaaccaag ccgcttcttg ggatgatgta cctgaaaaga ttaaagaaac ctttgacgt 360
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gaagtgggtt accacaacat gaagggaag ttccaaaaat taggtattat ctttacagat 480
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ccgcgcagac ataacaagtt ggcagccctc aactcagag tatggtcggg tggaactttt 600
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tctgaggcaa ctgaaatgat tgtcatggga ttgtagaac cctttacaaa agaacttcca	1380
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taa	1443

<210> 492

<211> 1254

<212> DNA

<213> Streptococcus pneumoniae

<400> 492

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gcaaatactg gtaaaattct ctatgagaag gatgcaacgc aacctgtcga aattgcttcc	180
ataacaaaat tgattactgt ttatctggtc tatgaagctt tggaaaacgg aagtattacc	240
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aaaaaatacc cacaagtctt agaaatcacc aaaaaacctt cttctacttt tgctgggatg	660

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gctggaacag tggttggcca ttgacttat gaagacaagg acttgattgg tcaaggttac	1140
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ttcttaaaag tttggtggaa tcagtttgtc cgctttgtta acgagaaatt ataa	1254

<210> 493

<211> 1074

<212> DNA

<213> *Streptococcus pneumoniae*

<400> 493

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gccatccgtg cagttgttcg tcaagcaatt tcagaaggaa tggaaagttt tggatatctat	180
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 atccaattcg ccaaggaaagc tggttgctcc ctccaaggga cggaagatcc agaggccaat 1440
 catatcaacg ttattaagag aaagacaaac tataagaaaa gtcnatag 1488

<210> 496
 <211> 861
 <212> DNA
 <213> Streptococcus pneumoniae

<400> 496
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<210> 497
 <211> 885
 <212> DNA

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 ggtactgcag aagaaggggc attgtttagc cttactgcag aaggttaagat tgtggttaac 1020
 aaccacaca aagccgatat tgagctatct agcttgaata agagcttgct ataa 1074

<210> 494
 <211> 1545
 <212> DNA
 <213> *Streptococcus pneumoniae*

<400> 494
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<210> 495
 <211> 1488
 <212> DNA
 <213> Streptococcus pneumoniae

<400> 495
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 aatccagaac ttgtcgaaact cttaggagaa aaatgtgtag gcattgatgt caatctgatg 180
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 gcaggggctg atatggcagc agtttccatg cataagtctg gtgggagttt gacccaaagc 720
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 acccagtccta catctgcctc ttacttgttg atggctagt ttgatatttc acgtcgcaac 840
 ttggcccttc gtggtaaga gtcgtttgag aaagtcattg agctatctga gtatgccgc 900
 cgtgaaatca atgctatcgg tggctactat gcctactcaa aagagttaat agacgggtgt 960
 tcggttttgc attttgacgt aactaagctg tcagtttaca ctacgggtat tggcttaaca 1020
 ggtatcgagg tttatgacct ctgtcgagac gaatacgaca ttcagatcga gtttgggtat 1080

<213> *Streptococcus pneumoniae*

<400> 497
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 attattctct tgcccagagt gtttgaacat cctattttct gtcaggaaacg tcaagtatgac 180
 tactaccagt atgcccaatc tgtagcggaa aatactgccca ttcagcattt taagggtgatt 240
 gctaaggaaac tacaagtgtt tttaccaatc agtttctatg aaaaagatgg taatgtcttg 300
 tataactcta ttgccgtcat tgatgcagat ggggaagtgc tgggcgttta tcgaaagacc 360
 catataccag atgaccatta ttatcaagaa aaattctatt tcacgcctgg taacactggt 420
 ttcaaggctc ggaatactcg ctatgctaag attggtatcg gtatctgttg ggatcaatgg 480
 ttccctgaaa cagcgcgctg tcttgcatg aatggtgctg aattgctctt ttatcctaca 540
 gctatcgggt cagagccaat tttggataca gatagttgtg gtcactggca acgtactatg 600
 caagggcacg cagcagcgaa tattgttcca gtcacgcag ccaatcgta ttggtttagag 660
 gaggttactc ctagttagga aaatggcgga cagagctcca gtcttgactt ctacggttcc 720
 tcctttatga cggatgaaac aggaactatt ctagaacgag ctgaaagaca agaagaagct 780
 gttctgttag ctacttatga cctagacaag ggagcaagtg aacgcctaaa ctggggcgtg 840
 ttctgagata gaagaccaga aatgtataga caaattacag attag 885

<210> 498

<211> 363

<212> DNA

<213> *Streptococcus pneumoniae*

<400> 498
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 ttagcaaaag gttactatgg agctaaacac atcttgttcc gtactgcaaa agaacaagta 120
 atgaactctt actactatgc ataccgtgac cgtcgtcaga aaaaacgtga cttccgcaaa 180
 ttgtggatta ctogtatcaa tggcgagct cgtatgaacg gactttcata ctacaattg 240
 atgcattggt tgaattggc tgagatcgaa gttaacgta aaatgcttgc tgacttggct 300
 gttacgatg cagtagcttt cacagctctt gcagatgcag ctaaagcaaa acttggtaaa 360
 taa 363

<210> 499
 <211> 555
 <212> DNA
 <213> *Streptococcus pneumoniae*

 <400> 499
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 atattttaa ac tgggtccaa ga tccaggttga ttaccagaag gcaagggaa c tgcccaacgg 120
 gatttgattt tccacaatgg ggctgtctgt gtttttagcag taacggatga acaaaaaactt 180
 atcttgg tca agcagtaccg caaagctatc gaggtgtgtc cttacgaaat tccagccgga 240
 aaattggaag taggagaaaa cacagcccct gtggcagctg cccttcgtga attagaggaa 300
 gaaacagcct atacaggga attagaactc ttgtacgatt ttatttcagc tattggcttt 360
 tgtaatgaga agttaaact atatttagca agcgatttga caaaagtga aaatccgcgt 420
 ccgcaggatg aggatgaaac ctgtgaagtc cttgaagta gcttagaaga agcgaagaa 480
 ttaatccaat caggtcatat ctgtgatgcc aagacaatta tggctgttca gtattgggag 540
 ttgcagaaaa aatag 555

 <210> 500
 <211> 585
 <212> DNA
 <213> *Streptococcus pneumoniae*

 <400> 500
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 caacagcaa aacaacaac gactgtacaa caaattgctg ttggaaaaa tgctccagac 180
 ttcacattgc aatccatgga tggcaagaa gttaagtatt ctgattttaa gggtaaaaag 240
 gtttacttga agttttgggc ttcattggtgt ggtccatgca agaaaagtat gccagagttg 300
 atggaaactag cggogaaacc agatcgtgat ttcgaaatc ttactgtcat tgcaccagga 360
 attcaaggty aaaaaactgt tgagcaattc ccacaatggt tccaggaaca aggatataag 420
 gatatccag ttctttatga taccaaagca accaccttc aagcttatca aattcgaagc 480
 attcctacag aatatttaat tgatagccaa ggaaagattg gaaagattca atttggtgct 540
 atcagtaatg cggatgcaga agcagcattt aaagaaatga actag 585

 <210> 501

<211> 1269
 <212> DNA
 <213> *Streptococcus pneumoniae*

<400> 501
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 aatgctattg ccaagaaga agaacgcaa caaataata tcgagttaat tgcttcggaa 120
 aacgtagttt ccaaggctgt tatggcagct caagggctta tcttgacaaa taaatatgcc 180
 gagggttacc caggagccgc ttattatggt ggaactgatg tagtagacgt tgtagagact 240
 cttgctattg aacgcgcaa agaaattttc ggtgctaaat ttgccaatgt tcaaccacat 300
 tcaggaagcc aagctaactg tcgggcttac atgtccttga ttgagccagg tgatacgggt 360
 atgggaatgg atttgccatc aggtgggtcat ttgactcatg gggctcctgt tagctctctc 420
 ggtcaaacct acaactttgt ttcttatagt gttgatccta aaacggaact cttagacttt 480
 gatgctatct tgaacaagc ccaagaagta aaacaaaac tgattgtagc tgggtgctca 540
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 aagctcatgg tggacatggc ccatatcgct ggcttgggtg cggctggcct tcatccaagc 660
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 ggtggttga ttttgaccaa tgacgaagaa cttgctaaaa aaatcaattc agctattttc 780
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 gaagtttttg atccagcctt caaggaatat gctgccaatg taattaagaa cagcaaggct 900
 atggcagatg tcttcttgca agacctgat ttccgtatta tttcagtggt aactgaaaac 960
 catctcttcc ttgttgatgt gactaaagtt gtgaaaaacg gcaaagttgc tcaaaacttg 1020
 ttggatgaag tcaatattac cttaataaaa aactcaatcc cttacgaaag cttgtcacca 1080
 ttcaagacaa gtgggattcg tatcggagca gcagccatta ctgcacgtgg atttgggtgaa 1140
 gaagaaagtc gcaaagtggc tgaactcatc attaaaacc ttaagaattc agaaaatgag 1200
 gctgtattag aagaagttag aagtgcagtc aaagaattga cagatgcctt cccattatc 1260
 gaggactaa 1269

<210> 502
 <211> 1035
 <212> DNA
 <213> *Streptococcus pneumoniae*

<400> 502
 ggagttttta tgaaaaaacaa attttttcta atagctattt tagctatgtg tatagttttt 60
 agcgcttggt cttctaattc tgttaaaaaat gaagaaaata cttctaaaga gcatgcgcct 120
 gataaaatag ttttagatca tgccttcggt caaactatat tagataaaaa acctgaaaga 180
 gttgcaacta ttgcttgggg aaatcatgat gtacgattag ctttaggaat agttcctgtt 240
 ggattttcaa aagcaaatga cgggtgaagt gctgataaag gagttttacc atggacagaa 300
 gaaaaaatca aagaactaaa tggtaaaagt aacctattt acgattttgga tggacttaac 360
 tttgaagcaa tatcaaatc taaaccagat gttatcttag caggttatc tgggtataact 420
 aaagaagatt atgacactct atcaaaaaatt gctcctgtag cagcatatac atctaaacct 480
 tggcaaacct tatggagaga tatgattaaa attgattcaa aagccttagg tatggaaaaa 540
 gaagtgtagt agttaataca aaatactgaa gctcgtatat ccaagaatt agaaaaacat 600
 ccagaaatca aaggaaaaat caaaggaaaa aaagtattt ttactatgat taatgctgca 660
 gatacatcaa aattctggat ttatactagc aaagatccaa gagcaaatca tttacagat 720
 ttaggcttag ttttcctga atcattaaaa gaatttgaga gtgaagatag ttttgcaag 780
 gaaatttctg cagaagaagc aaataagata aatgatgctg atgtaatcat aacttatggt 840
 gatgataaaa ctcttgaagc ttacaaaaa gatcctcttt taggtaaaaa aaatgcaatt 900
 aaaaaatggt cgttgctgtt aattccagat aatacacctg tagcagcctc atgcaactca 960
 acaccactt caataaacta tactattgaa gaatacctaa atcttttagg aaatgcagtc 1020
 aaaaatgcga aataa 1035

<210> 503

<211> 2700

<212> DNA

<213> *Streptococcus pneumoniae*

<400> 503

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 gaaaaataa tacagttgaa agaattatca acgcaggaag attatcaagg tctaaacctg 180
 ctagtgacta gcttatcaaa tgaatgaagt gtctatatct cagcgtattt ctctatcttg 240
 cctcttttga ttaatatctc agaggatgtg gatttagctt atgaaatcaa tcatcaaat 300
 aatattgac aggactattt aggtaaatta tctacaacga ttaaatgggt agcagaaaa 360

gaaaatgcgcg ttgagatcct agaacacttg aatgttgtgc ctgttttgac agcccatcca	420
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aaataccgtg atgttaagtt ggggttgatc aataaagata aatggtacca tgatttgcgt	540
cgttacatcg aaattatcat gcagacagac atgattcgtg agaaaaaatt aaaagtgact	600
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aataaagatg gcggttactt gtcacatgtt tggacctctc acaaggctca acaacaattg	1740
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cagaagaaga gcgataccaa taccocaaat cgttatgaaa ccattatgga tcaagtagtg 2040
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 ttccaatcgc ttctttcaaa tgttgatatg gttttgtcaa aatcaaatat gaattattgt 2400
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 aatgaatggc aagtacttaa gaacgttatc ttggctattg aaggacatga cgaactctta 2520
 gctgacaatc catatctaaa agctagtctg gattaccgta tgccttactt taattatttc 2580
 aactatatc agttggagtt gattaacgc caacgtcgtg gagaattgtc cagtgtatca 2640
 gaacgattga ttcatatcac catcaacgga attgcgacag gattgcgtaa ttcaggttga 2700

<210> 504
 <211> 372
 <212> DNA
 <213> Streptococcus pneumoniae

<400> 504
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 cgttacgatt acactcttag cccttctgtt aaaaaattca ccctcaaaga taacaccttt 120
 tttgaaaacta aggttggtaa ctatgaactg actgcctttt tggaaaaagt gccaaacagc 180
 ggtgaaggct tccaactcaa aatcatcatt aacaagggaac ttacaggggc taaaatcaat 240
 atcactgaca agtttggcct tcgtctagt gataatttca aatcagaaga ccaccatatt 300
 catcaggaaa aattctactt cctcatggat agcttggtag aacgtggtgt ctttacaaaa 360
 tcggaagat ag 372

<210> 505
 <211> 1050
 <212> DNA
 <213> Streptococcus pneumoniae

<400> 505
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 atttataaag ttacagaggy acggtctata atgtcagata gaaaaaacat gaaacttttc 120

gcactcaact ctaaccaaga gattgcacaa aaaattgcc aagctgttg tgcccactt 180
 ggaaaactat catcacgtca attttcagac ggagaaatcc aagtaaataat cgaagaaagt 240
 gtccgtgggt atgatgttta catcatccaa tcaacaagtt tccctgtcaa caaccaacta 300
 atggaattgt taatcatggt cgaatgtgt gtgcgtgcaa gtcccacag tatcaacgtt 360
 gtccctccat attttggtca tgcacgtcaa gaccgcatg ctgtccctcg tgagccactt 420
 acagcaaac tagttgccaa tatgctggtt aaggctggag ttgatcgtat cctgactctt 480
 gatttgcagt cgttcaggt tcaagggttc tttgatattc cagtggataa tcttttcaat 540
 gttccctat tgcacaaaca ttactgcgat aaaggattgc ttggttcaga tgttgttgtc 600
 gttagcccta aaaattcagg tgcacacgt gcgcgtagcc tggctgaata tcttgatgct 660
 cctatcgcca ttatcgacta cctcaagac gatgcaactc gtaacgaagg ttatattatt 720
 ggtgatgttg aagtaagaa agctatcttg attgatgata ttttaatac aggacgtacc 780
 ttctctgaag cttctaaat cgttgaacgt gaaggagcta cagaaattta tgctgtttct 840
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 attcttgtga ctgattcagt agcaacaaaa gaaaaaactc ctaaaaatgt atgctacatc 960
 actgctagt agttaattgg tgatgctatc gtccgtattc acgaaagaaa accagtcagc 1020
 ccactctttg cctacaataa aaagaaataa 1050

<210> 506

<211> 1470

<212> DNA

<213> *Streptococcus pneumoniae*

<400> 506

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 aatttagtaa atggaaatg gaaatcatct gaacaagaaa ttacgattta ttcaccaatc 120
 aatcaagaag aattgggtac agttccagcc atgactcaga ctgaagctga tgaggctatg 180
 caagctgcgc gtgcagccct gccagcatgg cgagctttat cagcagttga acgtgcggct 240
 tatttgcata aaacagcagc tatttttagaa cgcgataagg aagaaattgg tactatocct 300
 gccaaagaag tagcaaaagg gattaaagca gcaattggag aagtagtgcg tacagcagac 360
 ttgattcgtt atgctgctga ggaaggtctc cgtatcactg gacaagcaat ggaaggtggc 420
 ggttttagag caacaagtaa aaacaaactg gctgttgtcc gtgcgtgaacc agttggtatc 480

gtgctagcga ttgctccctt taattatcca gttaatttat ctgcttctaa aattgcacct 540
 gccttgattg cagggaatgt ggtcatgttt aagccacca cacaagggtc catttctgga 600
 ctcttggttg ctaaacgatt tgaagaagca gggattccgg cagggtgttt caacaccatt 660
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 gttacaaaag atatgaaagt ggcatgggaa gagccatttg gtcctgtttt accaatcatt 1200
 cgtgtggcta gtgtagagga agctattgcc tttccaacg aatctgaatt cggccttcaa 1260
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 ggtacagtc acattaataa taaacccag cgtggctcag ataatttccc attccttggt 1380
 gtcaaaggtt ctggagctgg agtgcaagga attaaatata gcattgaagc gatgacaaat 1440
 gtcaaatcca ttgtttttga tgtgaaataa 1470

<210> 507

<211> 993

<212> DNA

<213> *Streptococcus pneumoniae*

<400> 507

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 gttaaacacc aaacagaaga accaactgtg gcccaaatgg aagaacttaa agtcttggta 180
 gcagatgaat tgactaataa tgcttcactc atgcttcttg acctgagta tggacttcca 240
 gcaactaaag ctcttgatga aaaagctggg cttctccttg cttatgaaaa aacaggttat 300
 gacacacaa gcacaaaacg ctggccagac tgcttggatg ttgggtctgc aaaacgtatt 360
 aaagaagaag gtgcagatgc agttaaatc ttgctttact atgatgtaga tagctcagac 420


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gaactcaatc aagaaaaaca agcctacatc gaacgcatcg gttctgagtg tgtggctgaa 480
gatateccat tcttcttga aatccttgct tacgatgaaa aaattgcgga tgcaggttct 540
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ccacgcctta acattgatgt ttgaaagtt gaagttcctg ttaacattaa atatgttgaa 660
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ttccaagata ctcttgtatt tgctcatgaa tcaggtgcga actttaacgg agttcttctgt 840
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cgcgaatgcg ttgcgacaa cggatttgaa aacattgacg aactcaacaa agttcttcaa 960
agaacagcaa cttcatggaa agaacgcgtg taa 993

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<210> 508
<211> 546
<212> DNA
<213> Streptococcus pneumoniae

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<400> 508
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gactttggta cctatgacca tacacggact cactacccaa tctttggtaa aaaagtaggg 180
gaagctgtaa ctacgggtca agctgatctt ggagtatgta tctgtggtac tgggtgtggg 240
atcaacaacg ctgtaataaa agttccagggt gttcgttctg ccttggttcg tgatatgaca 300
acagcccttt atgctaaga acaattgaac gctaacgtta ttggtttttg tggtaaaatt 360
actggtgaat tgcttatgtg tgatatcatc gaagctttca tccatgctga atacaaacca 420
actgaagaaa acaaaaaatt gattgcgaaa attgaacatg ttgaaagtca caatgctcaa 480
caaacagacg caaacttctt tacagaattc cttgagaaat gggatcgtgg agaataccac 540
gactaa 546

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<210> 509
<211> 441
<212> DNA
<213> Streptococcus pneumoniae

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```

<400> 509

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```

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ggccaagact ttgttgatgt gactctcgtc gttgctgcag aagtaacaa agaagaacaa      180
aaccttggtat tcgtgattga tgcttatgga gctgggtccat ttatagttgc aactaagatc      240
aaaggaatgg ttgctgcaga agtatctgac gaacgttcag cttatatgac tcgtggccac      300
aacaactcac gtatgatcac tatggggagca caacttggtg gtgatgaatt ggcataaaat      360
atcgctaaag gatttgtaa tggtaaatac gacgggtggtc gtcaccaaata ccgggttgac      420
atgttgaaca aaatgggcta a                                     441

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<210> 510
<211> 1893
<212> DNA
<213> Streptococcus pneumoniae

```

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<400> 510
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attatcgccc atattgacca cggaaaaatca actctagcag accgcatttt ggaaaaaaca      180
gagacgggtt caagtcgtga aatgcaggcc cagcttttgg atagcatgga gctagagcgg      240
gaacgtggaa ttactattaa gttgaatgcc atcgagttga attacactgc aaaagatggg      300
gaaacttata tttccactt gattgacaca ccagggcacg ttgactttac ctatgaagtt      360
tcacgttcgc tagctgcctg tgaggggtgct attttgggtg tcgatgcggc tcaaggaatt      420
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cgtgtgggtg ataccgttac cttggcaacc aatcctgcgg cagaaccatt acatggttat      960

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 ttctcagcg tcttgagtat ggtgaagaa taa 1893

<210> 511

<211> 1347

<212> DNA

<213> *Streptococcus pneumoniae*

<400> 511

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 cgacctactc accaatattt ctccctaaag gatgaccatg cagttattca agcgaccatc 180
 tggctctgga tttatcagaa attagggttt gacctggaag aaggaatgaa gatcaatgtg 240
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<210> 512

<211> 750

<212> DNA

<213> Streptococcus pneumoniae

<400> 512

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aatattttca tagtctggaa taccaataag tatcaagtta gccaggtatc aaagaaaaa 180
ttagaagaaa atcaggatag agaaggcaat ttgactttg attctgtcaa agctatctct 240
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gctagtacc atactcttgg tgttgataat gctaataaaa tgttatcttc tcttttagat 480
aatgctaaaa atggcatgaa gatttatcta accgataaaa ataaagtta tacttatgaa 540

atacgtgaag tcaaacgtgt gacacccgat cgtgttgatg aagttgatga tagagatggg	600
gtcaatgaaa tcacattagt aacctgtgaa gaccttgctg ctacagaacg tattattgtc	660
aaaggtgatt tgaagaagaac aaaagattat tcacaacat ctaatgaaat cctaacagct	720
ttcaatcaac catataaaca attttattaa	750

<210> 513

<211> 1026

<212> DNA

<213> Streptococcus pneumoniae

<400> 513

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aaccaaggaa ttgcacaaga gcttggaatt atcgaaatcc cacaattgca tgaaaaagct	180
gttggtgatg cgcttgacct tagtcacgcc cttgccttca cttcaactaa aaaaatctat	240
gcagctcaat actctgactg tgcagacgct gaccttggtg tgatcactgc aggtgcacct	300
caaaaaccag gtgaaactcg tcttgacctt gtaggtaaaa accttgctat caacaaatca	360
atcgtaactc aagttgttga atctgggttc aaaggtatct tccctgttgc tgcataccca	420
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gtttggtcac acgctaactc cgtgggtgta aacctgaag aattccttaa agacactcaa	660
aatgttcaag aagctgaatt gattgaattg ttccgaaggtg ttctgtatgc agcctacaca	720
atcatcaaca aaaaagggtgc aacatactac ggtatcgcag tagcccttgc tcgtatcact	780
aaagcaatcc ttgacgatga aaacgcagta cttccacttt cagtattcca agaaggtaa	840
tacggagttg agaattgtct tatcgggtcaa ccagctgttg ttggtgcaca tgggtatcgtt	900
cgtccagtaa atatcccatg gaacgacgca gaaactcaaa aaatgcaagc atctgcataa	960
gaattacaag ctatcattga cgaagcatgg aaaaaccagg aattccaaga agcttctaaa	1020
aactaa	1026

<210> 514

<211> 588

<212> DNA

<213> *Streptococcus pneumoniae*

<400> 514
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 acgggttcaa tcgcctctta taagtcggca gatttagtca gttctctaaa aaaacaaggc 120
 catcaagtca ctgtcttaat gactcaggct gctacagagt ttatccaacc ttgacacta 180
 caggctactct cacagaatcc tgtccacttg gatgtcatga aggaacccta tctgtatcag 240
 gtcaatcata tcgaacttgg aaaaaagca gatttattta tcgtgggtacc tgcaactgct 300
 aacactattg caaaactagc tcacggattt gcggacaaca tggtaccag tacagctcta 360
 gccctacca gtcataatcc caaactaata gctcctgcta tgaatacaaa aatgtatgac 420
 catccagtaa ctcagaataa tctgaaaaa ttgaaacct acggctatca gctgattgct 480
 cctaaggaat cctactagc ttgtggagac cacggacgag gagctttagc tgacctcaca 540
 attattttag aaagaataaa ggaaactatc gatgaaaaa cgctctaa 588

<210> 515

<211> 924

<212> DNA

<213> *Streptococcus pneumoniae*

<400> 515
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 aaaatcatta aagaaaaaat ttcttcctta cttagtcaag aagagggaagt cctcagtggt 120
 gaacaactgg gtggaatgac caatcaaaac tatttggcca aaacaacaaa taagcaatac 180
 attgttaaat tctttggtaa agggacagaa aagcttatca atcgacaaga tgaaaagtac 240
 aatcttgaa ctaactaaagga tttaggctta gatgtaaaa attatctttt tgatattgaa 300
 gctggtatca aagtaaatga gtatatcgaa ctgcgatta cgcttgatc aacgtcaatc 360
 aagaccaagt tcgacaaaat tactccaata ttacaaacta ttcatacgtc tgctaaggaa 420
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tataaaattt tacaagatac tatttggagt ctatggactg tctataagga agagcaaggt 840
 gaagattttg gtgactatgg tgtgaatcgt taccaaagag ctattaaagg tttgcttct 900
 tatggagggt cagatgaaaa gtaa 924

<210> 516
 <211> 813
 <212> DNA
 <213> Streptococcus pneumoniae

<400> 516
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 gattatattg atactctctg taaaaagcac aatatcaact atattattaa ctacgggtact 120
 ctgattgggg cggttcgaca tgagggtctt atcccttggg acgacgatat tgatctgtcc 180
 atgcctagag aagactacca acgatttatt aacatttttc aaaaggaaaa aagcaagtat 240
 aagctcctat ccttagaagc tgataagaac tactttaaca actttatcaa gataaccgac 300
 agtacgacta aaattattga tactcgaat acaaaaacct atgagtcctg tatctttatc 360
 gatattttcc ctatagatcg ctttgatgat cctaagggtc ttgatacttg ttataaactg 420
 gaaagcttca aactgctgtc ttccagtaaa cataaaaata ttgtctataa ggatagcctt 480
 ttaaagatt ggatacgaac agccttctgg ttactccttc gaccggttcc tctcgttat 540
 ttbgcaata aaatcgagaa agaaattcaa aaatatagtc gtgaaaatgg gcaatatatg 600
 gcttttatcc cttaaaaatt taaggaaaag gaagtcttcc caagtgtgtac ctttgataaa 660
 acaatcgatt taccttttga gaatttaagc cttcctgcac ctgaaaaatt tgatactatt 720
 ttgacacaat tttatggaga ttatatgacc ctaccaccag aagaaaaacg cttctacagt 780
 catgaatttc acgcttataa attggaggat tag 813

<210> 517
 <211> 357
 <212> DNA
 <213> Streptococcus pneumoniae

<400> 517
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 ccacatttcc gtctcgttga cactgttcgt gtacatgcga aagttgtcga aggtaacgt 120
 gaacgtatcc agatttttga aggtgttgtt atcgacgta aaggtgtcgg aatctcagaa 180

aactacacag	ttcgtaaaaat	ctctaacggt	gtagggtgtg	agcgtatctt	cccaatccac	240
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<210> 518
 <211> 459
 <212> DNA
 <213> Streptococcus pneumoniae

<400> 518	
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gatgaattct	gtaaggctgt tgatgacttt gaccgtgtct ttgtgtcaac aggagcagaa 360
aaaggttcag	agtggtgtaa agttgatctt tctgccgagg aagaagatat tgaacgcttg 420
gaacaattcg	cagaagaatt ggctgctaaa gtaggataa 459

<210> 519
 <211> 438
 <212> DNA
 <213> Streptococcus pneumoniae

<400> 519	
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cttgaattga	acgacctgtg aaaagctatc gaagaagaat ttgggtgaac tgcagctgct 180
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actggtcttg	gtcttaaaaga agctaaagaa cttgttgacg gtgcaccagc acttgtttaa 360
gaaggcgttg	caactgcaga agctgaagaa atcaaagcta aattggaaga agctggagct 420
tcagttactc	ttaaataa 438

<210> 520
 <211> 627

<212> DNA
 <213> *Streptococcus pneumoniae*

 <400> 520
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 aaaaatgaaag ctgctgcacg tctcgtcgtt gtagacgctc gtggtttgac agtgagcaa 240
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 ggaccatctg cagtagcatt ttctaataaa gatgttatcg caccagcgaa aatcttgaac 420
 gacttttcta aaaacgctga agcacttgaa attaaagggt gtgcaatcga aggcgctgctc 480
 gcatctaaag aagagattct tgcacttgca actcttcaa accgcgaagg actctcttct 540
 atgctccttt ctgtacttca agcgccagtg cgcaacggtg ctcttgcagt caaagcggtt 600
 gcagaaagca aagaagacgc ggcttaa 627

 <210> 521
 <211> 1809
 <212> DNA
 <213> *Streptococcus pneumoniae*

 <400> 521
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 ctctcatga ctctcttagt ctgtgtgtg acgccaactg ccatgggtgat ttacagcttt 600
 attgccaaga aatcctatca tctcttccag aagcaaacag agacaggggg aattcagact 660

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cagttgattg aagaatcgct tagtcagcag actataatoc agtccttcaa tgctcaaaaca 720
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tctgaataa 1809

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<210> 522

<211> 1053

<212> DNA

<213> *Streptococcus pneumoniae*

<400> 522

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acaaaggaga aaaaaatcat gtttgaacat tattcagtag ctgatttgtt tgcaaatctt 60
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<210> 523

<211> 1329

<212> DNA

<213> *Streptococcus pneumoniae*

<400> 523

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agtttcatgc gtgataatgt ggtctatata gatggttata gtcaagtgat taatggtaga 660
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<210> 524

<211> 825

<212> DNA

<213> Streptococcus pneumoniae

<400> 524

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gatattgcta aagtaactgg gcagattctc tatcgtggaa ttgatgtcaa ccgtccagaa 300
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gctaagtc aa tttaccgtaa tattaccttt gcgcatgaac gtgctggagt taaggataag 420
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aaagcagtc tccacaagtc agccttgacc ttatcaggtg gtcagcaaca acgtctctgt 540
atcgctcgtg ccatctctgt taagccagat atcctcttaa tggatgagcc agcctcagcc 600
ttgatccga ttgcgaccat gcaactagaa gagaccatgt ttgagctcaa gaaaacttt 660

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accatcatca ttgtaacgca taatatgcag caggctgctc gtgcaagtga ctatacaggc	720
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atccacatgt tctcacttgc tatccaagag catgtgacaa ttgataaatt ggcattgaca 1320

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acggctgaaa aataa 1395

<210> 526
 <211> 1263
 <212> DNA
 <213> Streptococcus pneumoniae

<400> 526

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ggacacgttg accacggtaa aactacccta actgcagcta tcacaactgt tttggcacgt 180

cgcttgacct catcagttaa ccaacctaaa gactatgcgt ctatcgatgc tgctccagaa 240

gaacgcgaac gcggtatcac tatcaacact gcgcacgttg agtacgaaac tgaaaaacgt 300

cactacgctc acatcgacgc tccaggacac gcggactacg ttaaaaacat gatcactggt 360

gctgctcaaa tggacggagc tatccttgta gtactctcaa ctgacggacc aatgccacaa 420

actcgtgagc acatccttct ttcacgtcag gttggtgta aacaccttat cgcttctcatg 480

aacaagttg acttggttga cgacgaagaa ttgcttgaat tggttgaaat ggaaatccgt 540

gacctattgt cagaatacga cttcccaggt gacgatcttc cagtatatcca aggttcagca 600

cttaaagctc ttgaaggtga ctctaataac gaagacatcg ttatggaatt gatgaacaca 660

gttgatgagt atatccaga accagaaact gacactgaca aaccattgct tcttccagtc 720

gaggacgtat tctcaatcac tggacgtggt acagttgctt caggacgtat cgaccgtggt 780

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gttggtactg gtgttgaaat gttccgtaaa caacttgacg aaggtcttgc tggagataac 900

gtaggtgtcc ttcttcgtgg tgttcaacgt gatgaaatcg aacgtggaca agttatcgct 960

aaaccaggtt caatcaaccc acacactaaa ttcaaagggtg aagtctacat ccttactaaa 1020

gaagaagggtg gacgtcacac tccattcttc aacaactacc gtccacaatt ctacttccgt 1080

actactgacg ttacagggtc aatcgaaact ccagcaggta ctgaaatggt aatgcctggt 1140

gataacgtga caatcgacgt tgagttgatt caccacatcg ccgtagaaca aggtactaca 1200

ttctctatcc gtgaggggtg acgtactggt gggtcaggta tggttacaga aatcgaaact 1260

taa 1263

<210> 527
 <211> 873
 <212> DNA
 <213> *Streptococcus pneumoniae*

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 agtggtctga tgcctcatgc tgttggttat ggtatggggg ttatgatccc agccttgatg 180
 tttggtaacg tatctgggaa tcacatcaac cctgctttca ctctagggct tgcagtttagc 240
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<210> 528
 <211> 1416
 <212> DNA
 <213> *Streptococcus pneumoniae*

<400> 528
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 gaaagaaaaa caaaaatcgt ccttgaagta gccttggagt taggagatgg tatggttcgt 180
 actatcgcca tggaaatcaac agatgggttg actcgtggaa tggaaagtatt ggacacaggt 240
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 gaaatgaag aatcaggcgt tatcgagaaa acagccatgg tctttggtca gatgaatgag 660
 ccaccaggag cacgtatgcg tgttgccctt actggtttga caatcgctga atacttcgt 720
 gatgtggaag gccaaagcgt gcttctcttt atcgataata tcttcggtt cactcaggct 780
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 caagatatca ttgctatcct tggatggat gagctttctg atgaagaaaa gaccttggtt 1200
 gctcgcgccc tcgtatcca gttcttcttg tcacaaaact tcaacgttgc ggaacaattt 1260
 actggtcagc caggttctta tgttccagtt gctgaaactg tacgtggctt taaggaaatc 1320
 cttgatggta aatacgacca ctggccagaa gatgccttcc gtggtgtagg ttctatcgaa 1380
 gatgtgattg caaaagctga aaaaatggga ttttaa 1416

<210> 529

<211> 888

<212> DNA

<213> *Streptococcus pneumoniae*

<400> 529

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 gaagctgctc gcaacttcca agtttacgct cagaaagtc gtaaaacttt gagacatac 180
 cttcatggta atggagctgg tgcttcaact aatccgatgt tgattagccg ttctgtgaag 240
 aagacaggct atatcggttat cacttcagac cgcggtttgg ttggaggtta taattcctct 300


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at t t t t g a a a g   c t g t t a t g g a   g t t g a a a g a a   g a a t a c c a c c   c a g a c g g t a a   a g g t t t t g a a   360
a t g a t c t g t a   t c g g t g g g a t   g g g a g c t g a t   t t c t t t a a g g   c t c g c g g t a t   t c a a c c a c t t   420
t a t g a a t t a c   g t g g c t t g t c   a g a c c a a c c t   a g c t t t g a t c   a a g t t c g t a a   g a t t a t t t c a   480
a a a a c t g t t g   a a a t g t a c c a   a a a t g a a c t c   t t t g a t g a g c   t t t a t g t t t g   c t a c a a c c a c   540
c a t g t c a a t a   c g c t a a c c a g   t c a a a t c g c t   g t g g a a c a a a   t g c t t c c g a t   t g t t g a c t t g   600
g a t c c a a a t g   a a g c g g a t g a   a g a g t a c a g c   t t g a c t t t t g   a a t g g a a a c   c a g c c g a g a a   660
g a a a t t c t g g   a g c a g t t g t t   g c c t c a g t t t   g c a g a a a g t a   t g a t t a c g g   t g c c a t t a t c   720
g a t g c c a a g a   c a g c t g a g a a   t g c t g c g g g c   a t g a c a g c c a   t g c a a a c a g c   g a c a g a t a a t   780
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a c a c a a g a a a   t t a c a g a a a t   c g t a g c a g g t   g c t a g t g c c t   t a g a a t a g   888

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<210> 530

<211> 1536

<212> DNA

<213> *Streptococcus pneumoniae*

<400> 530

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t t a a t t a a g c   a a c a a a t t g a   a a a t t t c a a a   c c c a a t t t t g   a t g t g a c t g a   a a c a g g t g t t   120
g t a a c c t a t a   t c g g g g a c g g   t a t c g c g c g t   g c t c a c g g c c   t t g a a a a t g t   c a t g a g t g g a   180
g a g t t g t t g a   a t t t t g a a a a   c g g c t c t t a t   g g t a t g g c t c   a a a c t t g g a   g t c a a c a g a c   240
g t t g g t a t t a   t c a t c c t a g g   t g a c t t t a c a   g a t a t c c g t g   a a g g c g a t a c   a a t c c g c c g t   300
a c a g g g a a a a   t c a t g g a a g t   c c c t g t a g g t   g a a a g t c t g a   t t g g t c g t g t   t g t g g a t c c g   360
c t t g t c g t c   c a g t t g a c g g   t c t t g g a g a a   a t c c a c a c t g   a t a a a a c t c g   t c c a g t a g a a   420
g c a c c a g c t c   c t g g t g t t a t   g c a a c g t a a g   t c t g t t t c a g   a a c a t t g c a   a a c t g g t t t g   480
a a a g c a t t a g   a g c c c c t t g t   a c c g a t t g g t   c g t g g t c a a c   g t g a g t t g a t   t a t c g g t g a c   540
c g t c a g a c a g   g g a a a c a a c a   c a t t g c g a t t   g a t a c a a t c t   t g a a c c a a a a   a g a t c a a g a t   600
a t g a t c t g t a   t c t a c t c g c   g a t t g g a c a a   a a g a a t c a a   a g t t c g t a c   g c a a g t a g a a   660
a c a c t t c g t c   a g t a c g g t g c   c t t g g a c t a c   a c a a t c g t t g   t g a c a g c c t c   t g c t t c a c a a   720
c c a t c t c c a t   t g c t c t t c c t   a g c t c c t t a t   g c t g g g g t t g   c t a t g g c g g a   a g a a t t a t g   780
t a t c a a g g t a   a g c a t g t t t t   g a t t g t a t a c   g a t g a t c t a t   c a a a a c a a g c   g g t a g c t t a t   840

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<210> 531
 <211> 567
 <212> DNA
 <213> Streptococcus pneumoniae

<400> 531
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 tcagacttga ctcaaatcaa gcaagttgtt gaaaaaacag gtctgccttc ttttttaaaa 180
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 tatgatgtgc ttgtagattg cttgaaccga cttgaaaaag aaacaatcg atttgaagtg 360
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 gagaaaaaaa tgtctctgaa agtaaggagt gtaaaagaac aaatcgatga aagtcctatt 480
 ggtggttttg tcattttttg caatcacaag acaattgatg tgagtattaa acaacaactt 540
 aaagttgtta aagaaaattt gaaatag 567

<210> 532
 <211> 498

<212> DNA
 <213> *Streptococcus pneumoniae*

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 gaaagagctg aaaaaattgc ttcatgatat gacagagctg aagaagcccg tcaaaaagca 180
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 gaagctttac agagtgttaa ggggtagggtc gcagatttga ccacagctt agctggtaaa 420
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<210> 533
 <211> 1419
 <212> DNA
 <213> *Streptococcus pneumoniae*

<400> 533
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 ccaatttatc ccgacttaga ttagatgcc gatttccatg aaaaagaaga tatctatcag 480
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<210> 534
<211> 252
<212> DNA
<213> Streptococcus pneumoniae

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<400> 534
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ttogttttag aacgtgggaa aatccttctc cgtcgtgtaa caggaacttc agtcaaaaac    180
caacgtaaaag taacaacagc tatcaaacgc gctcgcgtaa tggctttgat gcctttcgta    240
aacgaagatt aa      252

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<210> 535
<211> 312
<212> DNA
<213> Streptococcus pneumoniae

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<400> 535
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ggtgcaactg ttgttgaatc aaaaacttgg gaaaaacgtc gtcttgcata cgaatccaa    180
gatttccgtg aaggacttta ccacatcggt aacgttgaag caaatgacga tgcagctctt    240
aaagagtttg accgtctttc aaaaatcaac gctgacattc ttcgtcacat gatcgtcaaa    300

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attgacgcgt aa 312

<210> 536
 <211> 1638
 <212> DNA
 <213> *Streptococcus pneumoniae*

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 tgggaatttc tcttcttctt ctgtgggcat ttgttagttg tggcaatatt atttggcttt 180
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 <213> Streptococcus pneumoniae

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<210> 538
 <211> 891
 <212> DNA
 <213> *Streptococcus pneumoniae*

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<210> 539
 <211> 1260
 <212> DNA
 <213> *Streptococcus pneumoniae*

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 gcttttattt gtaaatgaat cgtggagtta gctcaggaaa tcattcgaga agaattgggt 180
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<210> 540
 <211> 1164
 <212> DNA
 <213> *Streptococcus pneumoniae*

<400> 540
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 gataaagaat ttatcgtttt cgtaggacct tcaggatgtg gtaaatcaac tacactccgt 180
 atgattgctg gtcttgaaga cattacagaa ggtactgcat ctatcgatgg cgtagttgtc 240
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 ccacacatga ctgtttatga caacatggct ttcggtttga aatttcgtaa atacagcaaa 360
 gaagacatta acaaacgtgt tcaagaagca gctgaaatc ttggattgaa agaattcttg 420
 gaacgtaaac cagctgacct ttcagggtgt caacgtcaac gtgttgccat gggcgctgcg 480
 attgtccgtg atgcgaaggt attcttgatg gacgaacctt tgtcaaacctt ggatgccaaa 540

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gaaactgaaa aaacaatcta ctaa 1164

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<210> 541
<211> 1638
<212> DNA
<213> Streptococcus pneumoniae

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<400> 541
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gaacatgtca agattggggc taaggaattg acaacagaat tggttgacca gtactatata 720
cgtgttaagg aacaagaaaa atttgacacc atgactcgtc tcatggatgt ggcacaacca 780

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aacaaggcgc ataaataa 1638

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<210> 542
<211> 1317
<212> DNA
<213> Streptococcus pneumoniae

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<400> 542
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<210> 543

<211> 2370

<212> DNA

<213> *Streptococcus pneumoniae*

<400> 543

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<210> 544
 <211> 294
 <212> DNA
 <213> Streptococcus pneumoniae

<400> 544
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 acttgggaaa tcaaccacot taacgaacac atcaaaacaac acaaaaaaga ccacgtctact 180
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<210> 545
 <211> 1953
 <212> DNA
 <213> Streptococcus pneumoniae

<400> 545
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<210> 546

<211> 708

<212> DNA

<213> Streptococcus pneumoniae

<400> 546

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<210> 547
 <211> 549
 <212> DNA
 <213> *Streptococcus pneumoniae*

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 <211> 1389
 <212> DNA
 <213> *Streptococcus pneumoniae*

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<210> 549
 <211> 2073
 <212> DNA
 <213> Streptococcus pneumoniae

<400> 549
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<210> 550
 <211> 930
 <212> DNA
 <213> Streptococcus pneumoniae

<400> 550
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 gatgatatcg taaaagaaaa tatgattgaa actctttatc agcaagtcca agaaaaggat 360
 gcagatgttg ttatagggaa ttactataat tatgacgaaa gtgaacgggaa tttttatttt 420
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 gatgcagaac aaagaatggt taaagttga 930

<210> 551
 <211> 345
 <212> DNA
 <213> Streptococcus pneumoniae
 <400> 551

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tgctgtatgc aaggtccaat cttggacaaa ttgtctgaag aactttcaga agatgtcttg	180
aaaatcgtaaat aaatggacgt tgatgaaat ccaaacacag ctctgtcttt tggaatcatg	240
tctattccaa ctctctctct caaaaaagac ggccaagttg tcaacaagt tgcaagtggt	300
cacacagcag aacaatacaa ggccatcatt gctgaattga gctaa	345

<210> 552

<211> 624

<212> DNA

<213> Streptococcus pneumoniae

<400> 552

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aaaatcattg gtctttcact agaaaacgtt tcaggtcttt tgggaatcga tgggtggttc	180
ttctcaaatc ttaaagaaaa aatcgttaac agcgatgacg taacaagtgg tgttaacgta	240
gaagttggta aaacacaaagt tgcagttgac ttaaacgtta ttggtgagta ccaaaaaaat	300
gttcagcgtt tatattcaga aatcagagaa atcgtatctt cagaagttgc taaaatgact	360
gacttggaat ttgttgaaat caacgtaaac gttgtcgaca tcaaaactaa agaacagcat	420
gaagcagact cagtaagcct tcaagatcgc gtatctgacg ttgctgaatc aacaggagaa	480
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cacgaaaaca ctctgtgtaa ctaa	624

<210> 553

<211> 1764

<212> DNA

<213> Streptococcus pneumoniae

<400> 553

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gtcattaaaa gtgtcatacc cctcgtggct toccacttta tcgaccagta tctcagcaat	180
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tctgggattt tatccagctt tatctcagca gtttttatct ttctgacaac cctttatacc 480
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1764

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<210> 554

<211> 480
 <212> DNA
 <213> *Streptococcus pneumoniae*

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 agtatagagg ctgaaaaatc agatgaagcc gaacgacttg taagagaaaa gtataagagt 420
 tgtgaaattg ttcttgatgc agatgatttt caggactatg acactagcat atatgaatag 480

<210> 555
 <211> 1014
 <212> DNA
 <213> *Streptococcus pneumoniae*

<400> 555
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 gaagatgatg gtttccgct ttcagttact gttttacgct atggttctat ctaccgtttg 240
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gaacctaagc gtgaaaaatgg ctttggtctat gatccctctc tcctttaggc agaaacaggc 900
 gagtcacacg ctgaattaac cctggaagaa aaaaatagtc aatctcaccg tgccttagcc 960
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<210> 556
 <211> 318
 <212> DNA
 <213> Streptococcus pneumoniae

<400> 556
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 gctctaggag gaatgtactt ggttcgtaag caaatcgaaa aagaatttgc tgacaacca 180
 cgtttgaatg ctgaagcagt tcgtactctt ttgagtcaa atgggtcaaaa accaagcgaa 240
 gctaaggtac aacaagtta ccaccaaatc atccgccaac aaaaggcagc cttgtctaac 300
 aataaaaaga aaaaataa 318

<210> 557
 <211> 312
 <212> DNA
 <213> Streptococcus pneumoniae

<400> 557
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 ttggttgcct caagtgttaa aactggagat cgtgtcttag ttgaagccca cgcaggctct 240
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 attgaggaat ag 312

<210> 558
 <211> 477
 <212> DNA
 <213> Streptococcus pneumoniae

<400> 558
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agtatccagt	caaaagaagt	ggcagctagt	gcagatgctc	cagttaaaaa	agagtag	477
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<212> DNA						
<213> Streptococcus pneumoniae						
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<210> 560

<211> 3720

<212> DNA

<213> Streptococcus pneumoniae

<400> 560

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<210> 561

<211> 1011

<212> DNA

<213> *Streptococcus pneumoniae*

<400> 561

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<210> 562
<211> 837
<212> DNA
<213> Streptococcus pneumoniae

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<210> 563
<211> 1224
<212> DNA
<213> Streptococcus pneumoniae

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<400> 563
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<210> 564

<211> 528

<212> DNA

<213> Streptococcus pneumoniae

<400> 564

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aaggaaagcg  acttggtctt  tgtcaaggga  gttcattctg  ctctacgaaa  cgacgtataa  180
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gggtccaacg cctttgaacg cctcttggtc ggctcttcac ctgaatacat actccgccat 480
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<210> 565
<211> 1113
<212> DNA
<213> Streptococcus pneumoniae

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<210> 566
 <211> 591
 <212> DNA
 <213> *Streptococcus pneumoniae*

<400> 566
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 tttgatttcg atgttgagat tgggtcaact gtacgtatta ttgatggtgc ttttgcagac 480
 tacactggta agattacaga aattgataat aacaaagtga aatgattat ctctatgttt 540
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<210> 567
 <211> 2205
 <212> DNA
 <213> *Streptococcus pneumoniae*

<400> 567
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<210> 568
 <211> 1221
 <212> DNA
 <213> *Streptococcus pneumoniae*

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 aaggttcgta aatacgggtc tcattggtaca agtcaccagt ttgtagcagg agaagctgca 600
 aaactcttgg gacgtccatt agaagacttg aagttaatta cctgtcatat tggtaacgga 660
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 ggttgatgat ttgatgatga aaagaatgct tttggcgtaa caggagacat ctcaacagag 1140
 gcagctaaaa tccgtgtcct gggtattcca acagatgaag aattagtcac tgcccgtgac 1200
 gttgaacgct tgaaaaaata a 1221

 <210> 569
 <211> 1098
 <212> DNA
 <213> *Streptococcus pneumoniae*

 <400> 569

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actataaata ttacaaataa aaaaacggag gagtgcttta tgaagccta tacttatggt      60
aaaccaggac ttgcttcttt tgttgatgta gacaaaccag ttattcgcaa gccaacagac      120
gctattgtgc gtattgtaaa aaccactatt tgtggaacag acctccatat tatcaaaggg      180
gattgttcta cttgccaaag tggtagcatt cttggccacg aagggtattg gattgttgaa      240
gaagtgtggg aaggagtctt caacttcaa aaagggtgaca aggtcttgat ttcttgcgct      300
tgtgcctgtg gtaaatgcta ctactgtaa aaagggaatt atgctcactg tgaagcga      360
gggggctgga ttttcggta cttgattgat ggtatgcagg ctgaatatct acgtgtccct      420
catgcagata atactcttta ccatactcca gaagacttgt catatgaagc tttggttatg      480
ctgtcagaca ttctgcctac tggatatgaa attggtgtct taaaggga agtagaacct      540
ggtgtcagcg tagccattat tggttcaggt ccagttggat tggctgtctt ttaacagcc      600
caattctatt caccagctaa attgattatg gtagacctag acgataaccg cttggaaact      660
gccctatcat tcggtgcgac tcataaggtt aattcttcag accctgaaaa agccattaaa      720
gaaatttatg atttgacaga tggtcgtggt gtggatgtcg ctatcgaagc tgttggtatt      780
cctgcaacat ttgatttctg tcaaaagatt atcggtgtag acggaacggt tgccaactgt      840
ggtgtgcatg gtaaacacgt tgaattcgat ttagataaac tttgattcgc caacatcaat      900
gtaacaactg gtttggtatc tacaatacgc actccacaat tgttgaaagc acttgaaagt      960
cataagattg aaccgaaaaa attggttaact cactatttca aactcagtga aattgaaaaa      1020
gcctacgaag tcttcagtaa ggcagcagac caccatgcca ttaagggtcat tatcgaaaac      1080
gatattctcag aagcctaa                                     1098

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<210> 570
<211> 1704
<212> DNA
<213> Streptococcus pneumoniae

```

```

<400> 570
ggagaaagta tgctgattca gaaaataaaa acctacaagt ggcaggccct ggcttcgctc      60
ctgatgcagc gcttgatggt tgctagtcca cttctgcaac cgcgttatct gcaggaaagt      120
ttaggcgccc tccttactgg gaaatatgaa gctatttata gtatcggggc ttggttgatt      180
ggtgtggccg tagtcggtct agtgctggtt ggactcaatg ttgtcctcgc agcctatatt      240
gccaagggag ttctatccga ccttcgggag gatgccttcc gtaaaattca aaccttttct      300

```

```

tatgctgata ttgaacaatt taatgcggga aatctagtcg ttccaatgac aaatgatatc 360
aaccagattc agaacgttgt catgatgacc ttccaaattc ttttcagact tccctctttg 420
ttcatcggtt cgtttatcct agcgggtcaa accttaacct ctctgtggtg ggtgattgtt 480
ctcatggtag tcttgatttt tggtttgact gctgtcatga tgggaatgat ggggcctcgt 540
tttgccaagt ttcaaacctt tcttgagcgc atcaatgcca ttgccaagga aaatttacgt 600
ggcgttcgtg tgggtcaagtc ctttgtccaa gaaaaagagc aatttgctaa gtttacagag 660
gtctcagacg agcttcttgg tcaaaacctt tacattgggt atgccttttc agtagtggaa 720
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ggacgattga ttggtcaagg tacgcatgca gacttggttg ccaacaatgc cgtttaccgt 1680
gaaatctatg aaacacagaa atga 1704

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<210> 571
<211> 1050
<212> DNA
<213> Streptococcus pneumoniae

<400> 571

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aataagagaa agaactctat caaggaggaa atcatggaaa aacaaccgt cgcgctcttg      60
gggcctgggt cttgggggaa cgcctcttca caagtcttaa atgacaatgg acacggaggta    120
cgtatttggg gaaatcttcc cgagcaaatc aatgaaatta atacacacca tactaataag      180
cactacttta aagatgtcgt tctagacgaa aatatcattg cctacaccga cttagacagaa    240
acattgaaag atgtggatgc gattttgttt gttgtcccaa caaagtgac acgacttgtt      300
gccagcaag ttgcacaaac cttggaccat aaggttatca tcatgcacgc atcaaaggga      360
ttagaacctg atagccataa acgattatca accattcttg aagaagaaat tcctgaacat      420
ctccgtagt atctgtcgtg tgtttcaggg cctagtcatt cagaagagac cattgtgcgt      480
gacctactt taataactgc tgcctctaaa gatttcaaaa cagctcaata cgttcagaag      540
ctatttagta atcactactt ccgactttat accaatacgg atgttatcgg ggttgaaact      600
gctggtgctc ttaaaaaat tattgctgct ggtgctggag cttacatagg tcttggtatt      660
ggtgataatg ctaaggcagc catcatcgct cgaggtttag cagaaatcac ccgcctaggg      720
gtagcactcg gggccagtc attgacctat agcggcttat ctggtgtggg agatttgatc      780
gtaacgggaa cttccatcca ctctcgtaac tggagagctg gagatgctct cggaagagga      840
gaatccctag ctgatataga agctaatat ggcatggtaa tcgaaggaat ttcaaagact      900
cgagcagcct atgaactagc ccaagaactt ggagtcata tgcaccattac acaggctatt      960
taccaagtta ttatcacgg aaccaatata aaagatgcaa ttatgacat catgaacaat    1020
gaatttaaag cagaaaatga gtggtcttaa      1050

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```

<210> 572
<211> 1284
<212> DNA
<213> Streptococcus pneumoniae

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<400> 572
gaggaaataa ctatgtcatc taaatttatg aagagcgtg cgggtgcttg aactgctaca      60
cttgctagct tgccttttgg agcttgcgga agcaaaactg ctgataagcg tctgtattct      120
ggttcactcg aagtc aaaga actcaactgta tatgtagacg agggatataa gagctatatt      180
gaagagggtg ctaaaagctta tgaaaaagaa gctggagtaa aagtcactct taaaactggg      240
gatgctctag gaggtettga taaactttct cttgacaacc aatctggtaa tgcctctgat      300
gttatgatgg ctccatacga ccgtgtaggt agccttggtt ctgacggaca actttcagaa      360

```

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gtgaaattga gcgatggtgc taaaacagac gacacaacta aatctcttgt aacagctgct 420
aatggtaaag ttacggtgc tcctgccgtt atcgagtcac ttgttatgta ctacaacaaa 480
gacttggtga aagatgctcc aaaacattt gctgacttgg aaaccttgc taaagatagc 540
aaatacgcat tcgtggtga agatggtaaa actactgcct tcctagctga ctggacaaac 600
ttctactata catatggact tcttgccgtt aacggtgctt acgtcttgg ccaaaacggt 660
aaagacgcta aagacatcgg tcttgcaaac gacggttcta tcgtaggat caactacgct 720
aaatcttgtt acgaaaaatg gcctaaaggt atgcaagata cagaagggtc tggaaactta 780
atccaaactc aattccaaga aggtaaaaca gctgctatca tcgacggacc ttggaagct 840
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ggaaagaat atgctgcatt cgttggtggt aaagcttggg tcattcctca agccgttaag 960
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aaaaacgatg agttgacaac agctgttatc aaacagttca agaactca accactgcc 1140
aacatctctc aaatgtctgc agtttgggat ccacgaaaa atatgctctt tgatgctgta 1200
agtggtaaaa aagatgctaa aacagctgct aacgatgctg taacattgat caaagaaaa 1260
atcaaaaaa aatttggtga ataa 1284

```

```

<210> 573
<211> 1023
<212> DNA
<213> Streptococcus pneumoniae

```

```

<400> 573
caggaggta tcttatgac tgaaaaacta atcaattcaa aaccaaatgg tgtattcgca 60
ttgatctca ttgagttgac aatcgtaact ggtatcttta tatttataat gggggttgg 120
tcggaaaaca tttttggaat tattatcgga ctttactaa tcgtaattgc aggtctagct 180
catgctggtt taaaagttgt caaacctcaa gaagctctgg ttctgacact ctttggttaac 240
tatacaggta ccatcaaaaga acctggcttt tactttgtca atcccttcag cgtagcagtc 300
aacctgcaa accacactcg acttgacaaa agtggtgatg ttgacacaaa atctcctttt 360
ttaggagcta aatcatcaaa tgacaatgat gtaaatcttg aaattggcaa gaaacagatt 420
tccctcaaag tcattgacct gagcaattct cgtcaaaaa tcaatgattg cttaggaac 480

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cctgtagaaa tcggtatcgc ggtaacttgg agagttgtcg ataccgctaa ggcagtccttc 540
aacgttgata actacaaga atatctttca ttgcaatgtg atagcgccct ccgtaatat 600
gtccgcatct atccttacga tgtgtctcct aatgtggata ctacgggtga tgggcaagca 660
gatgaaggtg gtctccgtgg ctctagcgaa attgttgcta accgtattcg tgaagaaatc 720
caaagtcgtg ttgaggatgc tggcttgaa atccttgaag cagctatcac ttacctagct 780
tatgtccag aaattgtgc cgttatgctt caacgccaac aagcatctgc cattattgat 840
gcacggaaga tgattttaga tgggtctgta ggaatggtg aaatggcact agaacggctc 900
aatgaagggg aattggtaga acttgacgaa gaacgaaaag ctgccatggt ttcaaatctc 960
cttgtcgttc tttgtggcaa tcatgatgca caaccaattg tcaacacagg aagcctttac 1020
taa 1023

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```

<210> 574
<211> 1041
<212> DNA
<213> Streptococcus pneumoniae

```

```

<400> 574
aaaatagaaa gaggaataa taaatgaca aattcagtat tccaaggacg cagcttctta 60
gcagaaaag actttaccgc tgcagagtta gaatacctta ttggtcttcc agctcacttg 120
aaagatttga aaaaacgcaa tattcaacac cactaccttg ctggcaagaa tatcgctctc 180
ctatttgaaa aaacatctac tcgtactcgt gcagccttta caactgctgc tatcgacctt 240
ggtgctcacc cagaatacct cggagcaaat gatattcagt tgggtaaaaa agaattctact 300
gaagatactg ctaagattt gggacgtatg ttgacggga ttgaattccg cggattcagc 360
caacgtatgg ttgaagaatt ggcagaattc tcaggcgctc cagtattgaa cggctctaact 420
gacgaatggc acccaactca aatgctcgtc gactacttga ctgttcaaga aaacttcggt 480
cgcttggaag gcttgacatt ggtatactgt ggtgatggac gtaacaacgt tgccaacagc 540
ttgctcgtaa caggtgctat ccttggtgtc aatgttcaca tctctcacc aaaagaactc 600
ttcccagaaa aagaatcgt tgaattggca gaaggatttg ctaaagaag tggcgcacat 660
gttctcatca ctgaagatgc tgatgaagca gttaaagatg cagacgttct ttacacagac 720
gtttgggtat caatgggtga agaagacaaa ttcgcagaac gtgtagctct tcttaaacct 780
taccaagtca atatggactt agttaaaaa gcaggcaatg aaaacttgat ctctctacac 840

```

tgcttgccag cattccacga tactcacact gtttatggta aagacgttgc tgaaaaattt 900
 ggtgtagaag aaatggaagt aacagacgaa gtcttccgca gcaagtaacg tcgccacttc 960
 gatcaagcag aaaaccggtat gcacactatc aaagctgtta tggctgtctac acttggtaac 1020
 ctttatattc ctaaagtata a 1041

<210> 575
 <211> 924
 <212> DNA
 <213> Streptococcus pneumoniae

<400> 575
 aaaaataactt gtggagggttc cattatggca atatttttca tgatttttct gattgtttgt 60
 gtgctectat tgggtgatagt cacactgagt acagtttatg tggttcgta cgcgtcgggtg 120
 gcgattattg aacgcttttg gaaataccaa aaggttgcta atagcggat tcatattcgc 180
 ttgccttttg ggattgactc gattgcagca cggattcagt tgcgcttggc gcaaatgat 240
 attgtggttg agactaagac caagacaat gtgttcgtta tgatgaatgt agcgactcag 300
 taccgtgtca acgagcagag cgtgacagat gcttactata aactcatacg tccagaatct 360
 cagattaaat cttatatacga agatgctctt cgctcttctg ttccaaaatt aaccttggat 420
 gaattgtttg agaaaaaaga tgagattgcc cttgaagttc aacaccaagt agcagaagaa 480
 atgaccaact acggctacat tatcgtgaaa accttgatta ccaaggtcga accagatgca 540
 gaagttaagc aatccatgaa tgaaatcaat gcggcgcaac gtaagcgggt cgcagacaaa 600
 gaattggcgg aagctgacaa gattaaaatt gtcactgcag ctgaagcaga agcagaaaaa 660
 gaccgccttc atggtgtggg gattgcccga caacgtaagg cgattgtgga tggattggca 720
 gagtctatca ccgaactcaa ggaagccaat gttggcatga cagaagaaca aatcatgtct 780
 atcctcttga ccaaccagta ttggataacc ttgaatacct ttgcctctaa aggaatctaa 840
 actatctttt taccaaatat tccaaatggt gtggatgata tccgaacaca aatcttgcga 900
 gcccttcgcg ctgagaagaa ataa 924

<210> 576
 <211> 558
 <212> DNA
 <213> Streptococcus pneumoniae

<400> 576
 agattatcct ttaccagcgc cctttttctt ttaaaaatga gaaaatttcg gtataatagt 60

```

caaacacagggt caagggttttaa agagagagaggt ggggtttgtta tgagattttaa aaatacatcg 120
gatcatattg aggcctacat caaggcgatt ttagatcaat ctgggtatcgt ggagttgcaa 180
cggagtcagt tggcagatag ctttcagggt gttccatgac agatttaacta cgtgatcaag 240
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attcgtatag gacggattga gttttctagt catcatgaaa tgctccggga gctgctttac 360
tcgattggtg agcgagtcag tcaagaaatt tatgaggata ttctccagct tttggttgag 420
caggaattga tgaccaagca ggagatgaat ttgctagaat cagtagcttt ggatcgcggt 480
ttaggagaag aagctccagt tgttcgagca aacatgctac gtcagatcat acaagaggta 540
gatagaaaag ggaagtaa 558

```

<210> 577

<211> 477

<212> DNA

<213> Streptococcus pneumoniae

<400> 577

```

aatgaaggaa aaggagaaag aagaatgaaa gtaatctttt tagcagatgt taaagaaaa 60
ggtaaaaaag gcgaaattaa ggaagtacca acagggtagt cgcaaaactt tcttatcaaa 120
aagaatctag ccaaagaagc gactgctcaa gctgtagggt aacttcgtgg taacaaaaaa 180
tcggaagaaa aagctcacgc tgagatgatt gcagaaggaa aagcaattaa agcacactt 240
gaagcagaag aaactgttgt agaatttgtt gaaaaagttg gtccagatgg tcgtacctt 300
ggttctatta ccaataagaa gattgcagaa gaattgcaa agcaatttgg aattaagatt 360
gataaacgtc atattcaagt acaagctcgc attcgagcgg ttggtttgat tgatgtgcc 420
gtgaaaatct atcaagatat cacaagtgtg atcaatcttc gtgtgaaag aggataa 477

```

<210> 578

<211> 564

<212> DNA

<213> Streptococcus pneumoniae

<400> 578

```

aaagaaggta ctcatatgat taaatatagt atccgtggtg aaaacctaga agtaacagaa 60
gcaattctgt attatgtagt ttctaaactc gaaagatcg aaggtactt ccaaccagaa 120
caagagttgg atgcccgat taacttaaaa gtttatcgtg aaaaacggc taaagtgaa 180

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gtaacgattc cgettggtac tattactctc cgcgcagaag atgtatctca agatatgtat 240
 ggttcaattg accttgtaac tgataaaatt gaacgtcaga ttctgtaaaa taaaacaaaa 300
 atcagagcgt aaaaataaaa taaggtagca actggtcaat tatttacaga tgctttggtg 360
 gaagattcaa atattgtcca gtctaaagtt gttogttcaa aacaaattga tttaaaacca 420
 atggatttgg aagaagcaat tctacaatg gatttattgg ggcattgatt cttattctat 480
 gtggatgttg aagatcagac aaccaatgtg atttatcgtc gtgaggatgg cgaaattggt 540
 ttgttagagg ttaagaatc ttaa 564

<210> 579
 <211> 1080
 <212> DNA
 <213> Streptococcus pneumoniae

<400> 579
 cccgctctc ctattttcaa aaaatatag agaattaaaa tggcagaat tacagctaaa 60
 cttgtaaaag agttgctga aaaatctggt gccggtgta tggacgctaa aaaagcgctt 120
 gtagaaacag acggtgacat cgaaaaagcg attgaattgc ttctgtaaaa aggtatggct 180
 aaggcagcta agaaagctga ccgtgttgct gcagaaggtt tgactggtgt ttatgttaac 240
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 caattcgttg aattggttaa tactacagct aaagtcattg ctgaaggaaa acctgctaac 360
 aacgaagaag ctcttctttt gataatgcct tcaggtgaaa ctcttgaaac tgcatacgt 420
 tctgcaacag caactatcgg agagaaaatc tcattccgtc gctttgcatt gattgaaaa 480
 acagacgcac aacactttgg agcatacca cataacggtg gacgtatcgg tgttatttca 540
 gttgttgaag gtggagacga agcacttgct aaacaattgt caatgcacat cgcagcgatg 600
 aaaccaacag ttctttctta caaagaattg gatgagcaat tcgttaaga tgagttggca 660
 caattgaatc acgttatcga ccaagacaac gaaagccgtg caatggtaa taaacagct 720
 cttccacact tgaagtattg atcaaaagct caattaactg atgatgttat tgctcaagct 780
 gaagtgcac tcaaaagctga attggtgcga gaaggcaaac cagaaaaaat ctgggacaaa 840
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 acacttcttg cacaagttta catcatggat gacagcaaga cagttgaagc ataccttgaa 960
 tcagttaacg ctctggtagt tgagtttgcg cgtttgaag ttggtgaagg tatcgagaaa 1020

gctgcaaacg acttcaagc tgaagttgca gctacaatgg cagcagcctt gaataactaa 1080

<210> 580

<211> 873

<212> DNA

<213> *Streptococcus pneumoniae*

<400> 580

tactcattct tgaatgaattg tgaagcagtt gcccttgggt cgttttgcga gttgaagtca 60

agaagaggaa aaaacaaaa aggagaaata ctcatggcag taatttcaat gaaacaactt 120

cttgaggctg gtgtacactt tggtcaccaa actcgtcgtt ggaatcctaa gatggctaag 180

tacatcttta ctgaacgtaa cggaaatccac gttatcgact tgcaacaaac tgtaaaatac 240

gctgaccaag catacagactt catgcgtgat gcagcagcta acgatgcagt tgtattgttc 300

gttggtacta agaaacaagc agctgatgca gttgctgaag aagcagtacy ttcagggtcaa 360

tacttcacaa accaccgttg gttgggtgga actcttaca actgggggaa aatccaaaaa 420

cgtatcgttc gtttgaaga aattaaacgt atggaagaag atggaacttt cgaagttctt 480

cctaagaag aagttgcact tcttaacaaa caacgtgcgc gtcttgaaa atctctgggc 540

ggtatcgaag atatgcctcg tatcccgat gtgatgtac tagttgacc acataaagag 600

caaatcgtct ttaagaagc taaaaaattt ggaatcccag ttgtagcgat ggttgacacc 660

aatactgac cagatgatat cgatgtaac atcccagcta acgatgcgc tatcctgtct 720

gttaaatga toacagctaa attggctgac gctattatcg aaggacgtca aggtgaggat 780

gcagtagcag ttgaagcaga atttgcagct ttagaaactc aagcagattc aattgaagaa 840

atcgttgaag ttgtagaagg cgacaacgct taa 873

<210> 581

<211> 834

<212> DNA

<213> *Streptococcus pneumoniae*

<400> 581

tataaaggta gggatatgaa cggtttttaa aaatcaaaat atgtcattat tgtttttgtc 60

actgtttctc ttgtgtcagc tctcttagcg acgacttatt caagtacaa tgtgacaaaa 120

ttaggagatg gaatctcatt ggttgataga gttgtacaaa aaccttttca tgggtttgat 180

tctgtcaaat cagatttggc tcatttgaca cgaacatata atgaaaatga aagtttgaag 240

aaacagcttt accaattaga agttaaatca aatgaggtgg aaagtttaa gagagaaaa 300

gaacaactgc gccaatgtct tgatatgaag tctaaattgc aagccacaaa gacttttagca 360
 gcagatgtta ttatgcgttc tccggtatct tgggaagcagg agttgacctt agatgcagggt 420
 agatcaaaag gtgcttctga gaacatgtta gctattgcaa atggtggcctt gattgggaggt 480
 gtttcaaaag tagaggagaa ctctactata gtcaaccttc tgacaaatag ggaanaatgct 540
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 tatgacaagg aaaatgacgt tctttaaatt agccaattaa atagtaatag cgatattagt 660
 gcgggagata aggtgactac tgggtggatta ggaacttta acgttgctga tattcctgtt 720
 ggtgaagtgg ttgccacaac gcatagtaca gactatttga cagcagaagt aactgttataa 780
 ttgagtgcag atactcataa tgtagatgtg atagaattag tggggaattc ataa 834

<210> 582

<211> 846

<212> DNA

<213> *Streptococcus pneumoniae*

<400> 582

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 aaagaaaatc agaactacta cgatgtgaag gatattacgt ttcacgtgaa acgtggagaa 120
 tggctttcga ttgtagggca taatggtagt ggtaaatcaa cgacgggttcg attaatgtat 180
 ggcttacttg aagcagaatc cggagagatt gtaattgatg gccaacgttt gactaggagaa 240
 aatgtttgga atatacgtcg tcaaatcggt atggtttttc aaaatccaga caatcaattt 300
 gttggagoga ctgttgaaga tgatgttgct tttggttttg aaaatcaggg actttctcgt 360
 caagaaatga aaaagagagt ggaagaagct ctggcttttag ttggcatgtt ggactttataa 420
 aagagagagc cagcgcgtct atcaggtggc caaaagcaac gtgtggccat tgcagggtgtt 480
 gtagccctaa gaccagctat tttaatctta gatgaagcaa cgagtatgtt ggatcctgag 540
 gggcgtagag aactatttgg gacagtaaaa ggaattcgaa aagactatga tatgacggtc 600
 atttctatta cccatgattt ggaagaagtc gccatgagtg atcgcgtatt ggtcatgaaa 660
 aaaggggaaa ttgaatcaac tagtagtcca agggagcttt tctctcgaaa tgatttagat 720
 caaattggat tagacgatcc ttttgctaat caattaaaaa aatctttgag ccagaatggc 780
 tatgatttac ctgaaaatta tttagacaga agtgagctag aggataagct atgggaattg 840
 ctctag 846

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<210> 583
<211> 840
<212> DNA
<213> Streptococcus pneumoniae

<400> 583
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ttggaagcaa tgggaagcaga cgatttcgat caacttcaa gtctttttta caccgcttct      180
ttcttgaaaa aatatgcgatg ggctgttgag ttagatgacc aaattgtttt ggatgcttat      240
gattctggga gtatgattac ttataggaa gtagatgttg atgaagatga gttgacaggt      300
cgtagacgtt caagtaagaa aaagaagaaa aaaacatcat tttaccttt attttatttt      360
atcttttttg ctttatcgat tttaattttt gtgacttatt atgtttggaa ctatattcaa      420
actcaaccag aggagccttc tctttctaata tacagtgttg ttcaatcaac aagtccaact      480
agctctgttc cccactcctc aagtagtagt tcttctagta tagaatcagc tataagtgtta      540
tcaggcgaag gaaatcatgt agaaatcgct tataagacaa gtaaggaaac agttaaatgt      600
caattggcag tttcagatgt tacaagtgtg gtcagtgttt cagaaagcga acttgagggc      660
ggtgttaacct tatcgccaaa gaagaaaagt gcagaagcaa cagttgcaac taaaagtcct      720
gtaacaatta cgttaggtgt tgtaaaaggt gttgatttga cagtagataa tcagactgtt      780
gattttatcga aattaacagc tcagactgga caaatcactg taacctttac taaaaattaa      840

<210> 584
<211> 1020
<212> DNA
<213> Streptococcus pneumoniae

<400> 584
atggaaaatc ttgtgaaaag cacaagttat acatatatac cggaggaaat catgtctttt      60
tctgatttaa agctgtttgc ctttcttctt aataaagaat tggcagaacg tgggtgcgag      120
gagattggga tagagttggg gaaatcaagt gttcgccaat tttcagatgg agagattcag      180
gtcaacattg aagaatcaat ccgtgggaaa cagctcttta tcttacaatc aactagtctg      240
cctgtaaatg acaatctgct tgaaattttg attatggtag atgctttgaa cgcgtcgagt      300
gcagaatctg tcaatgttgt catgccttac tatgggtatg cagctcagga tagaaaggcg      360

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agagcgcgtg agccaatcac ttcaaaactt gtcgcaaata tgcttgaagt agctggagtg 420
 gatcgtttat tgaccatcga ctctcatgct ggcgcaaatc aaggattcct tgatattcct 480
 gtggatcatt tgatgggtgc tcctctgatt gcagattatt ttgagcgtcg tggatagggt 540
 ggttctgact atgtgggttg cagcccgagc catggagggg tgactcgtgc tcgtaagttg 600
 gcagaatttt tgaaaaacac tatcgctatt attgataaac gtcgtagcgt tgataagatg 660
 aatactagtg aagttatgaa tatcatcggt aaggttgaag gcaagacttg tatcttgatt 720
 gatgatatga ttgataccgc tggaaacgatt tgcctatgcg cagatgctct tgcggaagct 780
 ggtgctgttg aagcttatgc aagctgtacg caccagttc tttctggtcc tgctacggac 840
 aatatccaaa aatcagctat taagaaattg gttgttttgg ataccatcta tctgcagaaa 900
 gagcgtttga ttgataagat tgagcagatt tcaatcgctc atctcctagg ggatgctatc 960
 gtacgtattc atgaaaaacg accactttct ccacttttct atattgagaa aaaaatttaa 1020

<210> 585
 <211> 672
 <212> DNA
 <213> *Streptococcus pneumoniae*

<400> 585
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 ttgattcgtc agtttgacca ggctatttgc gagattcctg gggctcttgc tttgaccttg 120
 ggggaacctg attttacaac gccagaccat gtcaaggagg cgggcaagcg agcgattgat 180
 cagaaccaat cctactatac agggatgagt ggtctgctga ctctacgtca ggcagccagt 240
 gactttgtta aggaaaagta ccaactggac tatgctcctg aaaatgaaat cttggtttaca 300
 attggggcga cagaggtctt atctgcgact ttgacggcta ttttggaga gggagacaag 360
 gtacttttgc cagctcctgc ttatccaggc tatgaaccga ttgttaactt agttggggca 420
 gaaattgttg agattgatac gactgaaaaat ggttttgtct tgactcctga gatgttggag 480
 aaggccattt tggagcaggg tgataagctc aaggcgggta ttctcaacta tcagccaat 540
 ccgacaggaa ttacctacag tcgagagcag ttagaggcct tggcagctgt tttacgcaag 600
 tacgaaaatt ttgtgtctg tgatgaggtt tactcagaat tgacctacac agggogaagcc 660
 atgtgtctct ag 672

<210> 586

<211> 1302
 <212> DNA
 <213> *Streptococcus pneumoniae*

<400> 586
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 atccctaagg aagatgtggc ttgtattcgc aagaaggcgg actttgacat cgaccgtatt 180
 ttggaaattg agcaggagac gcgccacgat gtgggtgctt tcacgcgtgc ggtttctgag 240
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 actgcttatg gttacctcta caagcaggcc aacgacatca tcogtcgtga ccttgaaaac 360
 ttcactaata tcactgctga caagccaag gacacaaagt tcaccatcat gatggggcgt 420
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 gaaatgaaac gcaatatcga gcgcttcgag catgcggctg ctggtgtaga agtgtgtaag 540
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 gataaacttg gcatccgtgc ccaagaatc tctacacaag tccttctctg tgacctctac 660
 gctgagtact ttgcggttct tgccagcatt gcgacttcaa tcgaacgtat ggcgactgag 720
 attcgtggtc tacaaaaatc tgagcaacgc gaagtagaag agttctttgc taaggggcaa 780
 aaagggtctt cagcaatgcc tcacaaacgc aaccaatcg gttctgaaaa tatgactggt 840
 ctggcgctg tcattctgtg tcacatgatt acggcttatg aaaacgtcgc tctctggcat 900
 gaacgcgata tttctcactc atcagctgag cgtatcatca caccagatac gaccattttg 960
 attgactaga tgctcaacgc ttttggaat atcgtcaaga acttgacagt ctcccgaga 1020
 aatatgatcc gaaacatgaa ctgcactttt ggtcttatct ttacccaagg ggtatgttg 1080
 acattgattg aaaaaggcat gaccctgag caagcctatg acttggtgca accaaaaaca 1140
 gctactctt gggacaacca agtagacttt aaaccacttc ttgaggcaga ttcagaagta 1200
 acatcacgtc tcacacaaga agaaatcgat gaaatcttca acccagttta ttacacaaa 1260
 cgagtggtg atattcttga acgtcttgga ctagggtgatt aa 1302

<210> 587
 <211> 918
 <212> DNA
 <213> *Streptococcus pneumoniae*

<400> 587
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attactgggt taatcatggg agatgtgact actggtttac ttatcgggtg taacttgcaa 180
ctgtctgttc ttgggggttg taccttcggg ggtgcttctc gtatcgacgc aacttctggt 240
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acaatcgctg taccagtagc agctctcttg acttacttcg acgttcttg tgatgatgact 360
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gaacgcaact acttgcttgg tgcgattccg tgggctctat ctctgacctc tccagctctc 480
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aatgggttg cagatggctt gacacttgca ggacgtatgc ttccaggtct tggatttgca 600
atcttgcttc gttaccttcc agttaaacgt aaccttactc accttgctat gggatttggt 660
ttgacagcta tgttgactgt tctttactca tatgtaacag gtcttggtgg cgctggtgct 720
ggtatcgtag gtactcttcc tgctgaagtt gctgaaaaa ttggttctgt gaacaacttc 780
aaaggtttgt ctatgattgg tatttctatc gtaggatttt tccttgacgt gcttcaactc 840
aaaaatagcc aaaaagtagc tgtagcagca ccttctacac catcagaaag tggggaaatc 900
gaagatgacg aattctaa 918

<210> 588
<211> 855
<212> DNA
<213> *Streptococcus pneumoniae*

<400> 588
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tggaactacg aacgtatgca agcttctggt tacctttaca tgatcttgcc tcagttgogt 180
aaaatgtatg gtgatggaac tctgaattg aaagaaatga tgaaagtcca tactcaattc 240
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aaagatggtg taggttcaaa agacgccggt aacggtatca agacaggttt gatgggacca 360
ttcgctcctc ttggggatag aatctttggt tcacttctac ctgctatcat ggggtcagtc 420
gcagcaacta tggctatcgc tggccaacct tgggggatct tcctttggat tgcagttgca 480

gtagcgtatg acatcttccg ttggaacacg ttggaatttg cttacaaaga agggggttaac 540
 cttatcaaca acatgcaaag taccttgaca gctttgattg acgctgcac tgtacttggt 600
 gtcttcatga tgggtgctct tgtagcaaca gtgattaact ttgaaatttc ttacaagttg 660
 ccaatcggtg aaaagatgat tgatttccaa gacatcttga accaaatcct cccacgtttg 720
 cttccagcaa tctttactgc ctttatcttc tgggtgcttg gtaagaaagg tatgaactct 780
 actaaagcta tcggtattat tatcgtactt gctttggctc tttctgcctt tggctacatt 840
 gcacttgga tgtaa 855

<210> 589

<211> 1182

<212> DNA

<213> *Streptococcus pneumoniae*

<400> 589

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 caagcaaac gtgaagaat tgcagccttc ctacaagaaa tcgctgataa acatgactat 180
 attaaggta tcttgacagg tgctgggact tctgcttatg tgggagatac cttgctacct 240
 tattttaagg aagtctatga cgaacgcaa tgggaatttc atgctattgc gacaacagat 300
 atcgttgcca atccagcaac ctatttgaaa aaagatgtgg caactgtcct tgtgtctttt 360
 gctcgtagtg ggaattgcc tgaagtttg gcgactgttg atttggccaa atccttggtg 420
 gatgagcttt atcaagtgc gattacttgt gcagcagatg gtaaatggc tcttcaagct 480
 cacggtgatg atcgtaatct cttgctcttg caaccagctg tctctaata tctgtgattt 540
 gccatgactt cttagcttac gtctatgatg ttgacaactc tcttggctct tgatctcata 600
 gaatttgctg ttaagtctga acgttttgaa gttgtateta gtottgccc taaagtttta 660
 gacaaggcag aagatgtcaa agagctcgtt gatttagact ttaaccgtgt catctatcta 720
 ggcgtgtgtc ctttctttgg acttgctcat gaagctcagc tcaagatttt ggaattaaact 780
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gtcttgaatg atattttaccg tgtcttcctt tacatcggtt atgcccaact ctttgcttta 1080
 ttgacttcac tcaaggtaga aaataaacca gataccogt ctccacaggt tacagtaaac 1140
 cgtgtagtac aagggtgcat aattcacgaa tatcaaaagt aa 1182

<210> 590
 <211> 654
 <212> DNA
 <213> Streptococcus pneumoniae

<400> 590
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 ccactcttga aacaagctcg tcgtcttggc ctttcactta cagggtacaggt taaagaattg 120
 gcacgtcgta actacgtacc aggacaacac ggaccaaaaca accgttctaa attgtcagaa 180
 tacggtttgc aattggctga aaaacaaaaa cttcgtttca cttacggtgt aggtgaaaaa 240
 caattccgta acttggtctg acaagctaca aaaatcaaac gcggaatcct aggtttcaaac 300
 tttatgcttc ttttggaacg tcgtttggat aacgttggtt accgtccttg tctcgcgact 360
 actcgtcgtc aagctcgtca attcgtaac caccggtcaca tctctgttga cgggaaacgc 420
 gttgatatcc catcataccg cgtaactcca ggtcaagtga tctcagttcg tgaaaaaatca 480
 ttgaaagtcc cagcaatcct tgaagcagta gaagctactc ttggacgtcc agcattcgta 540
 tcattcgacg ctgaaaaaatt ggaaggttca ttgactogct tgccagaacg cgacgaaatc 600
 aaccagaaaa tcaacgaagc actgtgcgtt gaattctaca acaaaatgtt gtaa 654

<210> 591
 <211> 1032
 <212> DNA
 <213> Streptococcus pneumoniae

<400> 591
 ataaaaaatt tagaaaattt aagaatagaa aagagaacaa atcttatggc aaaagatatt 60
 cgtgtcttac ttactactct ttactatcca attgaaaatg cagagcaatt tgctgcgac 120
 cacttggtct tctgtaaatc aatcggtcctt aaaggccgta tctagtgcg tgacgaggga 180
 attaacggaa cagtttcagg tgactatgaa acaactcaaa aatacatgga ctacgttcac 240
 agccttcag gaattggaaga actctgggtc aagattgacg aagaaaatga acaagccttc 300
 aagaagatgt ttgttcgcta caagaaagaa attgtccacc ttggtttgga agacaacgac 360

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tttgacaatg acatcaaccc acttgaaaca acaggtgctt acttgtctcc aaaagagttc 420
aaagaagcgc ttcttgataa agataccggt gtccttgaca cagctaacga ttatgagtac 480
gacctaggac atttcctggt agctattcgc ccagatattc gcaacttcgc tgagttacca 540
caatgggtcc gtgataacaa ggaaaaattc atggacaagc gtgtcgtggt ttactgtaca 600
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ggccaattgc acggaggaat cgcaacttac ggtaagacc cagaagtcca aggtgagctt 720
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cgtggatgct cacacgagtg ccgtgttcac ccacgtaacc gctatgttcc aaaaaatgaa 960
ttgacacaag ctgaagttat cgagcgcccta gccgctatcg gtgaaagctt ggatcaagca 1020
gctactgtat aa 1032

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<210> 592
<211> 216
<212> PRT
<213> Streptococcus pneumoniae
<400> 592

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Met Asn Ile Ala Val Ile Gly Leu Gly His Val Gly Leu Ala Tyr Ala
1 5 10 15

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Leu Leu Phe Ala Ser Lys Tyr Lys Val Val Ala Tyr Asp Ile Asp Ser
20 25 30

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```

Val Lys Ile Asn Asn Leu Lys Lys Gly Ile Leu Pro Ser Lys Asn Glu
35 40 45

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Glu Leu Met Lys Phe Phe Cys Glu Asn Asn Leu Asn Ile Thr Phe Phe
50 55 60

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Asp Thr Phe Ser Glu Ile Lys Asn Asn Ile Asp Tyr Tyr Ile Ile Ala
65 70 75 80

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Leu Pro Thr Asp Tyr Asp Glu Lys Ile Gly Ser Phe Asn Thr Tyr Glu
85 90 95

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Ile Glu Gln Thr Val Ser Lys Ile Leu Arg Val Lys Pro Asn Gly Lys
 100 105 110

Ile Ile Leu Lys Ser Thr Val Pro Ile Gly Phe Ser Asn Lys Leu Lys
 115 120 125

Arg Leu Phe Asp Thr Lys Asn Ile Ile Phe Val Pro Glu Phe Leu Arg
 130 135 140

Glu Gly Cys Ser Ile Tyr Asp Asn Leu Tyr Pro Ser Arg Ile Val Val
 145 150 155 160

Gly Asp Glu Thr Val Glu Gly Arg Lys Ile Ala Glu Leu Phe Leu Ser
 165 170 175

Ile Ser Thr His Ser Thr Ala Asn Ile Lys Asn Val Met Leu Val Ser
 180 185 190

Pro Thr Glu Ala Glu Ala Ile Lys Leu Phe Ser Asn Thr Phe Leu Ala
 195 200 205

Leu Arg Val Ala Phe Leu Met Asn
 210 215

<210> 593
 <211> 234
 <212> PRT
 <213> Streptococcus pneumoniae

<400> 593

Met Val Leu Arg Gly Val Pro Met Ser Gln Ile Asp Leu Gln Lys Leu
 1 5 10 15

Thr Lys Lys Asn Gln Glu Phe Val His Ile Ala Thr Gln Gln Phe Ile
 20 25 30

Lys Asp Gly Lys Thr Asp Ala Glu Ile Gln Thr Ile Phe Glu Glu Val
 35 40 45

Ile Pro Gln Ile Leu Glu Glu Gln Ser Lys Gly Thr Thr Ala Arg Ser
 50 55 60

Leu Tyr Gly Ala Pro Thr His Trp Ala His Ser Phe Thr Val Lys Glu
65 70 75 80

Gln Tyr Glu Lys Glu His Pro Lys Glu Asn Asp Asp Pro Lys Leu Met
85 90 95

Ile Met Asp Ser Ala Leu Phe Ile Thr Ser Leu Phe Ala Leu Val Ser
100 105 110

Ala Leu Thr Thr Phe Phe Ala Ala Asp Gln Ala Phe Gly Tyr Gly Leu
115 120 125

Ile Thr Leu Leu Leu Val Gly Leu Val Gly Gly Phe Ala Phe Tyr Leu
130 135 140

Met Tyr Tyr Phe Val Tyr Gln Tyr Tyr Gly Pro Asp Met Asp Arg Ser
145 150 155 160

Gln Arg Pro Pro Phe Trp Lys Ser Val Leu Val Ile Leu Ala Ser Met
165 170 175

Phe Leu Trp Leu Leu Val Phe Phe Ala Thr Ser Phe Leu Pro Ala Ser
180 185 190

Leu Asn Pro Val Leu Asp Pro Leu Pro Leu Ala Ile Ile Gly Ala Ala
195 200 205

Leu Leu Ala Leu Arg Phe Tyr Leu Lys Lys Arg Leu Asn Ile Arg Ser
210 215 220

Ala Ser Ala Gly Pro Thr Arg Tyr Gln Glu
225 230

<210> 594

<211> 314

<212> PRT

<213> Streptococcus pneumoniae

<400> 594

Met Lys Gln Val Phe Leu Ser Thr Thr Thr Glu Phe Lys Glu Ile Asp
1 5 10 15

Thr Leu Glu Pro Gly Thr Trp Ile Asn Leu Val Asn Pro Thr Gln Asn
 20 25 30
 Glu Ser Leu Glu Ile Ala Asn Thr Phe Asp Ile Asp Ile Ala Asp Leu
 35 40 45
 Arg Ala Pro Leu Asp Ala Glu Glu Met Ser Arg Ile Thr Ile Glu Asp
 50 55 60
 Glu Tyr Thr Leu Ile Ile Val Asp Val Pro Val Thr Glu Glu Arg Asn
 65 70 75 80
 Asn Arg Thr Tyr Tyr Val Thr Ile Pro Leu Gly Ile Ile Ile Thr Glu
 85 90 95
 Glu Thr Ile Ile Thr Thr Cys Leu Glu Pro Leu Pro Val Leu Asp Val
 100 105 110
 Phe Ile Asn Arg Arg Leu Arg Asn Phe Tyr Thr Phe Met Arg Ser Arg
 115 120 125
 Phe Ile Phe Gln Ile Leu Tyr Arg Asn Ala Glu Leu Tyr Leu Thr Ala
 130 135 140
 Leu Arg Ser Ile Asp Arg Lys Ser Glu Gln Ile Glu Ser Gln Leu His
 145 150 155 160
 Gln Ser Thr Arg Asn Glu Glu Leu Ile Glu Leu Met Glu Leu Glu Lys
 165 170 175
 Thr Ile Val Tyr Phe Lys Ala Ser Leu Lys Thr Asn Glu Arg Val Ile
 180 185 190
 Lys Lys Leu Thr Ser Ser Thr Ser Asn Ile Lys Lys Tyr Leu Glu Asp
 195 200 205
 Glu Asp Leu Leu Glu Asp Thr Leu Ile Glu Thr Gln Gln Ala Ile Glu
 210 215 220
 Met Ala Asp Ile Tyr Gly Asn Val Leu His Ser Met Thr Glu Thr Phe
 225 230 235 240

Ala Ser Ile Ile Ser Asn Asn Gln Asn Asn Ile Met Lys Thr Leu Ala
245 250 255

Leu Val Thr Ile Val Met Ser Ile Pro Thr Met Val Phe Ser Ala Tyr
260 265 270

Gly Met Asn Phe Lys Asp Asn Glu Ile Pro Leu Asn Gly Glu Pro Asn
275 280 285

Ala Phe Trp Leu Ile Val Phe Ile Ala Phe Ala Met Ser Val Ser Leu
290 295 300

Thr Leu Tyr Leu Ile His Lys Lys Trp Phe
305 310

<210> 595

<211> 102

<212> PRT

<213> Streptococcus pneumoniae

<400> 595

Met Ala Asn Lys Lys Ile Arg Ile Arg Leu Lys Ala Tyr Glu His Arg
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20 25 30

Gly Ala Gln Val Ala Gly Pro Ile Pro Leu Pro Thr Glu Arg Ser Leu
35 40 45

Tyr Thr Ile Ile Arg Ala Thr His Lys Tyr Lys Asp Ser Arg Glu Gln
50 55 60

Phe Glu Met Arg Thr His Lys Arg Leu Ile Asp Ile Val Asn Pro Thr
65 70 75 80

Gln Lys Thr Val Asp Ala Leu Met Lys Leu Asp Leu Pro Ser Gly Val
85 90 95

Asn Val Glu Ile Lys Leu
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<210> 596

<211> 208

<212> PRT

<213> Streptococcus pneumoniae

<400> 596

Met Thr Lys Gly Ile Leu Gly Lys Lys Val Gly Met Thr Gln Ile Phe
 1 5 10 15

Thr Glu Ala Gly Glu Leu Ile Pro Val Thr Val Ile Glu Ala Thr Pro
 20 25 30

Asn Val Val Leu Gln Val Lys Thr Val Glu Thr Asp Gly Tyr Asn Ala
 35 40 45

Ile Gln Val Gly Phe Asp Asp Lys Arg Glu Val Leu Ser Asn Lys Pro
 50 55 60

Ala Lys Gly His Val Ala Lys Ala Asn Thr Ala Pro Lys Arg Phe Ile
 65 70 75 80

Arg Glu Phe Lys Asn Val Glu Gly Leu Glu Val Gly Ala Glu Ile Thr
 85 90 95

Val Glu Thr Phe Ala Ala Gly Asp Val Val Asp Val Thr Gly Thr Ser
 100 105 110

Lys Gly Lys Gly Phe Gln Gly Val Ile Lys Arg His Gly Gln Ser Arg
 115 120 125

Gly Pro Met Ala His Gly Ser Arg Tyr His Arg Arg Pro Gly Ser Met
 130 135 140

Gly Pro Val Ala Pro Asn Arg Val Phe Lys Gly Lys Asn Leu Ala Gly
 145 150 155 160

Arg Met Gly Gly Asp Arg Val Thr Ile Gln Asn Leu Glu Val Val Gln
 165 170 175

Val Val Pro Glu Lys Asn Val Ile Leu Ile Lys Gly Asn Val Pro Gly
 180 185 190

Ala Lys Lys Ser Leu Ile Thr Ile Lys Ser Ala Val Lys Ala Gly Lys
 195 200 205

<210> 597
 <211> 207
 <212> PRT
 <213> Streptococcus pneumoniae

<400> 597

Met Ala Asn Val Thr Leu Phe Asp Gln Thr Gly Lys Glu Ala Gly Gln
 1 5 10 15

Val Val Leu Ser Asp Ala Val Phe Gly Ile Glu Pro Asn Glu Ser Val
 20 25 30

Val Phe Asp Val Ile Ile Ser Gln Arg Ala Ser Leu Arg Gln Gly Thr
 35 40 45

His Ala Val Lys Asn Arg Ser Ala Val Ser Gly Gly Gly Arg Lys Pro
 50 55 60

Trp Arg Gln Lys Gly Thr Gly Arg Ala Arg Gln Gly Ser Ile Arg Ser
 65 70 75 80

Pro Gln Trp Arg Gly Gly Gly Val Val Phe Gly Pro Thr Pro Arg Ser
 85 90 95

Tyr Gly Tyr Lys Leu Pro Gln Lys Val Arg Arg Leu Ala Leu Lys Ser
 100 105 110

Val Tyr Ser Glu Lys Val Ala Glu Asn Lys Phe Val Ala Val Asp Ala
 115 120 125

Leu Ser Phe Thr Ala Pro Lys Thr Ala Glu Phe Ala Lys Val Leu Ala
 130 135 140

Ala Leu Ser Ile Asp Ser Lys Val Leu Val Ile Leu Glu Glu Gly Asn
 145 150 155 160

Glu Phe Ala Ala Leu Ser Ala Arg Asn Leu Pro Asn Val Lys Val Ala
 165 170 175

Thr Ala Thr Thr Ala Ser Val Leu Asp Ile Ala Asn Ser Asp Lys Leu
 180 185 190

Leu Val Thr Gln Ala Ala Ile Ser Lys Ile Glu Glu Val Leu Ala
 195 200 205

<210> 598
 <211> 98
 <212> PRT
 <213> Streptococcus pneumoniae
 <400> 598

Met Asn Leu Tyr Asp Val Ile Lys Lys Pro Val Ile Thr Glu Ser Ser
 1 5 10 15

Met Ala Gln Leu Glu Ala Gly Lys Tyr Val Phe Glu Val Asp Thr Arg
 20 25 30

Ala His Lys Leu Leu Ile Lys Gln Ala Val Glu Ala Ala Phe Glu Gly
 35 40 45

Val Lys Val Ala Asn Val Asn Thr Ile Asn Val Lys Pro Lys Ala Lys
 50 55 60

Arg Val Gly Arg Tyr Thr Gly Phe Thr Asn Lys Thr Lys Lys Ala Ile
 65 70 75 80

Ile Thr Leu Thr Ala Asp Ser Lys Ala Ile Glu Leu Phe Ala Ala Glu
 85 90 95

Ala Glu

<210> 599
 <211> 277
 <212> PRT
 <213> Streptococcus pneumoniae
 <400> 599

Met Gly Ile Arg Val Tyr Lys Pro Thr Thr Asn Gly Arg Arg Asn Met
 1 5 10 15

Thr Ser Leu Asp Phe Ala Glu Ile Thr Thr Ser Thr Pro Glu Lys Ser
 20 25 30

Leu Leu Val Ala Leu Lys Ser Lys Ala Gly Arg Asn Asn Asn Gly Arg
 35 40 45

Ile Thr Val Arg His Gln Gly Gly His Lys Arg Phe Tyr Arg Leu
 50 55 60

Val Asp Phe Lys Arg Asn Lys Asp Asn Val Glu Ala Val Val Lys Thr
 65 70 75 80

Ile Glu Tyr Asp Pro Asn Arg Ser Ala Asn Ile Ala Leu Val His Tyr
 85 90 95

Thr Asp Gly Val Lys Ala Tyr Ile Ile Ala Pro Lys Gly Leu Glu Val
 100 105 110

Gly Gln Arg Ile Val Ser Gly Pro Glu Ala Asp Ile Lys Val Gly Asn
 115 120 125

Ala Leu Pro Leu Ala Asn Ile Pro Val Gly Thr Leu Ile His Asn Ile
 130 135 140

Glu Leu Lys Pro Gly Arg Gly Gly Glu Leu Val Arg Ala Ala Gly Ala
 145 150 155 160

Ser Ala Gln Val Leu Gly Ser Glu Gly Lys Tyr Val Leu Val Arg Leu
 165 170 175

Gln Ser Gly Glu Val Arg Met Ile Leu Gly Thr Cys Arg Ala Thr Val
 180 185 190

Gly Val Val Gly Asn Glu Gln His Gly Leu Val Asn Leu Gly Lys Ala
 195 200 205

Gly Arg Ser Arg Trp Lys Gly Ile Arg Pro Thr Val Arg Gly Ser Val
 210 215 220

Met Asn Pro Asn Asp His Pro His Gly Gly Gly Glu Gly Lys Ala Pro
 225 230 235 240

Val Gly Arg Lys Ala Pro Ser Thr Pro Trp Gly Lys Pro Ala Leu Gly
 245 250 255

Leu Lys Thr Arg Asn Lys Lys Ala Lys Ser Asp Lys Leu Ile Val Arg
 260 265 270

Arg Arg Asn Glu Lys
 275

<210> 600
 <211> 93
 <212> PRT
 <213> Streptococcus pneumoniae

<400> 600

Met Gly Arg Ser Leu Lys Lys Gly Pro Phe Val Asp Glu His Leu Met
 1 5 10 15

Lys Lys Val Glu Ala Gln Ala Asn Asp Glu Lys Lys Lys Val Ile Lys
 20 25 30

Thr Trp Ser Arg Arg Ser Thr Ile Phe Pro Ser Phe Ile Gly Tyr Thr
 35 40 45

Ile Ala Val Tyr Asp Gly Arg Lys His Val Pro Val Tyr Ile Gln Glu
 50 55 60

Asp Met Val Gly His Lys Leu Gly Glu Phe Ala Pro Thr Arg Thr Tyr
 65 70 75 80

Lys Gly His Ala Ala Asp Asp Lys Lys Thr Arg Arg Lys
 85 90

<210> 601
 <211> 114
 <212> PRT
 <213> Streptococcus pneumoniae

<400> 601

Met Ala Glu Ile Thr Ser Ala Lys Ala Met Ala Arg Thr Val Arg Val
 1 5 10 15

Ser Pro Arg Lys Ser Arg Leu Val Leu Asp Asn Ile Arg Gly Lys Ser
 20 25 30

Val Ala Asp Ala Ile Ala Ile Leu Thr Phe Thr Pro Asn Lys Ala Ala
 35 40 45

Glu Ile Ile Leu Lys Val Leu Asn Ser Ala Val Ala Asn Ala Glu Asn
 50 55 60

Asn Phe Gly Leu Asp Lys Ala Asn Leu Val Val Ser Glu Ala Phe Ala
 65 70 75 80

Asn Glu Gly Pro Thr Met Lys Arg Phe Arg Pro Arg Ala Lys Gly Ser
 85 90 95

Ala Ser Pro Ile Asn Lys Arg Thr Ala His Ile Thr Val Ala Val Ala
 100 105 110

Glu Lys

<210> 602

<211> 217

<212> PRT

<213> Streptococcus pneumoniae

<400> 602

Met Gly Gln Lys Val His Pro Ile Gly Met Arg Val Gly Ile Ile Arg
 1 5 10 15

Asp Trp Asp Ala Lys Trp Tyr Ala Glu Lys Glu Tyr Ala Asp Tyr Leu
 20 25 30

His Glu Asp Leu Ala Ile Arg Lys Phe Val Gln Lys Glu Leu Ala Asp
 35 40 45

Ala Ala Val Ser Thr Ile Glu Ile Glu Arg Ala Val Asn Lys Val Asn
 50 55 60

Val Ser Leu His Thr Ala Lys Pro Gly Met Val Ile Gly Lys Gly Gly
 65 70 75 80

Ala Asn Val Asp Ala Leu Arg Ala Lys Leu Asn Lys Leu Thr Gly Lys
85 90 95

Gln Val His Ile Asn Ile Ile Glu Ile Lys Gln Pro Asp Leu Asp Ala
100 105 110

His Leu Val Gly Glu Gly Ile Ala Arg Gln Leu Glu Gln Arg Val Ala
115 120 125

Phe Arg Arg Ala Gln Lys Gln Ala Ile Gln Arg Ala Met Arg Ala Gly
130 135 140

Ala Lys Gly Ile Lys Thr Gln Val Ser Gly Arg Leu Asn Gly Ala Asp
145 150 155 160

Ile Ala Arg Ala Glu Gly Tyr Ser Glu Gly Thr Val Pro Leu His Thr
165 170 175

Leu Arg Ala Asp Ile Asp Tyr Ala Trp Glu Glu Ala Asp Thr Thr Tyr
180 185 190

Gly Lys Leu Gly Val Lys Val Trp Ile Tyr Arg Gly Glu Val Leu Pro
195 200 205

Ala Arg Lys Asn Thr Lys Gly Gly Lys
210 215

<210> 603

<211> 122

<212> PRT

<213> Streptococcus pneumoniae

<400> 603

Met Ile Gln Thr Glu Thr Arg Leu Lys Val Ala Asp Asn Ser Gly Ala
1 5 10 15

Arg Glu Ile Leu Thr Ile Lys Val Leu Gly Gly Ser Gly Arg Lys Phe
20 25 30

Ala Asn Ile Gly Asp Val Ile Val Ala Ser Val Lys Gln Ala Thr Pro
35 40 45

Gly Gly Ala Val Lys Lys Gly Asp Val Val Lys Ala Val Ile Val Arg
50 55 60

Thr Lys Ser Gly Ala Arg Arg Ala Asp Gly Ser Tyr Ile Lys Phe Asp
65 70 75 80

Glu Asn Ala Ala Val Ile Ile Arg Glu Asp Lys Thr Pro Arg Gly Thr
85 90 95

Arg Ile Phe Gly Pro Val Ala Arg Glu Leu Arg Glu Gly Gly Phe Met
100 105 110

Lys Ile Val Ser Leu Ala Pro Glu Val Leu
115 120

<210> 604

<211> 101

<212> PRT

<213> Streptococcus pneumoniae

<400> 604

Met Phe Val Lys Lys Gly Asp Lys Val Arg Val Ile Ala Gly Lys Asp
1 5 10 15

Lys Gly Thr Glu Ala Val Val Leu Thr Ala Leu Pro Lys Val Asn Lys
20 25 30

Val Ile Val Glu Gly Val Asn Ile Val Lys Lys His Gln Arg Pro Thr
35 40 45

Asn Glu Leu Pro Gln Gly Gly Ile Ile Glu Lys Glu Ala Ala Ile His
50 55 60

Val Ser Asn Val Gln Val Leu Asp Lys Asn Gly Val Ala Gly Arg Val
65 70 75 80

Gly Tyr Lys Phe Val Asp Gly Lys Lys Val Arg Tyr Asn Lys Lys Ser
85 90 95

Gly Glu Val Leu Asp
100

<210> 605
 <211> 180
 <212> PRT
 <213> Streptococcus pneumoniae
 <400> 605
 Met Ala Asn Arg Leu Lys Glu Lys Tyr Leu Asn Glu Val Val Pro Ala
 1 5 10 15
 Leu Thr Glu Gln Phe Asn Tyr Ser Ser Val Met Ala Val Pro Lys Val
 20 25 30
 Asp Lys Ile Val Leu Asn Met Gly Val Gly Glu Ala Val Ser Asn Ala
 35 40 45
 Lys Ser Leu Glu Lys Ala Ala Glu Glu Leu Ala Leu Ile Ser Gly Gln
 50 55 60
 Lys Pro Leu Ile Thr Lys Ala Lys Lys Ser Ile Ala Gly Phe Arg Leu
 65 70 75 80
 Arg Glu Gly Val Ala Ile Gly Ala Lys Val Thr Leu Arg Gly Glu Arg
 85 90 95
 Met Tyr Glu Phe Leu Asp Lys Leu Val Ser Val Ser Leu Pro Arg Val
 100 105 110
 Arg Asp Phe His Gly Val Pro Thr Lys Ser Phe Asp Gly Arg Gly Asn
 115 120 125
 Tyr Thr Leu Gly Val Lys Glu Gln Leu Ile Phe Pro Glu Ile Asn Phe
 130 135 140
 Asp Asp Val Asp Lys Thr Arg Gly Leu Asp Ile Val Ile Val Thr Thr
 145 150 155 160
 Ala Asn Thr Asp Glu Glu Ser Arg Ala Leu Leu Thr Gly Leu Gly Met
 165 170 175
 Pro Phe Ala Lys
 180

<210> 606
 <211> 132
 <212> PRT
 <213> Streptococcus pneumoniae

<400> 606

Met Val Met Thr Asp Pro Ile Ala Asp Phe Leu Thr Arg Ile Arg Asn
 1 5 10 15

Ala Asn Gln Ala Lys His Glu Val Leu Glu Val Pro Ala Ser Asn Ile
 20 25 30

Lys Lys Gly Ile Ala Glu Ile Leu Lys Arg Glu Gly Phe Val Lys Asn
 35 40 45

Val Glu Ile Ile Glu Asp Asp Lys Gln Gly Val Ile Arg Val Phe Leu
 50 55 60

Lys Tyr Gly Pro Asn Gly Glu Lys Val Ile Thr Asn Leu Lys Arg Val
 65 70 75 80

Ser Lys Pro Gly Leu Arg Val Tyr Lys Lys Arg Glu Asp Leu Pro Lys
 85 90 95

Val Leu Asn Gly Leu Gly Ile Ala Ile Leu Ser Thr Ser Glu Gly Leu
 100 105 110

Leu Thr Asp Lys Glu Ala Arg Gln Lys Asn Val Gly Gly Glu Val Ile
 115 120 125

Ala Tyr Val Trp
 130

<210> 607
 <211> 178
 <212> PRT
 <213> Streptococcus pneumoniae

<400> 607

Met Ser Arg Ile Gly Asn Lys Val Ile Val Leu Pro Ala Gly Val Glu
 1 5 10 15

Leu Ala Asn Asn Asp Asn Val Val Thr Val Lys Gly Ser Lys Gly Glu
 20 25 30

Leu Thr Arg Glu Phe Ser Lys Asp Ile Glu Ile Arg Val Glu Gly Thr
 35 40 45

Glu Ile Thr Leu His Arg Pro Asn Asp Ser Lys Glu Met Lys Thr Ile
 50 55 60

His Gly Thr Thr Arg Ala Leu Leu Asn Asn Met Val Val Gly Val Ser
 65 70 75 80

Glu Gly Phe Lys Lys Glu Leu Glu Met Arg Gly Val Gly Tyr Arg Ala
 85 90 95

Gln Leu Gln Gly Ser Lys Leu Val Leu Ala Val Gly Lys Ser His Pro
 100 105 110

Asp Glu Val Glu Ala Pro Glu Gly Ile Thr Phe Glu Leu Pro Asn Pro
 115 120 125

Thr Thr Ile Val Val Ser Gly Ile Ser Lys Glu Val Val Gly Gln Thr
 130 135 140

Ala Ala Tyr Val Arg Ser Leu Arg Ser Pro Glu Pro Tyr Lys Gly Lys
 145 150 155 160

Gly Ile Arg Tyr Val Gly Glu Phe Val Arg Arg Lys Glu Gly Lys Thr
 165 170 175

Gly Lys

<210> 608

<211> 118

<212> PRT

<213> Streptococcus pneumoniae

<400> 608

Met Ile Ser Lys Pro Asp Lys Asn Lys Leu Arg Gln Lys Arg His Arg
 1 5 10 15

Arg Val Arg Gly Lys Leu Ser Gly Thr Ala Asp Arg Pro Arg Leu Asn
 20 25 30

Val Phe Arg Ser Asn Thr Gly Ile Tyr Ala Gln Val Ile Asp Asp Val
 35 40 45

Ala Gly Val Thr Leu Ala Ser Ala Ser Thr Leu Asp Lys Glu Val Ser
 50 55 60

Lys Gly Thr Lys Thr Glu Gln Ala Val Ala Val Gly Lys Leu Val Ala
 65 70 75 80

Glu Arg Ala Asn Ala Lys Gly Ile Ser Glu Val Val Phe Asp Arg Gly
 85 90 95

Gly Tyr Leu Tyr His Gly Arg Val Lys Ala Leu Ala Asp Ala Ala Arg
 100 105 110

Glu Asn Gly Leu Lys Phe
 115

<210> 609

<211> 164

<212> PRT

<213> Streptococcus pneumoniae

<400> 609

Met Ala Phe Lys Asp Asn Ala Val Glu Leu Glu Glu Arg Val Val Ala
 1 5 10 15

Val Asn Arg Val Thr Lys Val Val Lys Gly Gly Arg Arg Leu Arg Phe
 20 25 30

Ala Ala Leu Val Val Val Gly Asp His Asn Gly Arg Val Gly Phe Gly
 35 40 45

Thr Gly Lys Ala Gln Glu Val Pro Glu Ala Ile Arg Lys Ala Val Asp
 50 55 60

Asp Ala Lys Lys Asn Leu Ile Glu Val Pro Met Val Gly Thr Thr Ile
 65 70 75 80

Pro His Glu Val Leu Ser Glu Phe Gly Gly Ala Lys Val Leu Leu Lys
 85 90 95

Pro Ala Val Glu Gly Ser Gly Val Ala Ala Gly Gly Ala Val Arg Ala
 100 105 110

Val Val Glu Leu Ala Gly Val Ala Asp Ile Thr Ser Lys Ser Leu Gly
 115 120 125

Ser Asn Thr Pro Ile Asn Ile Val Arg Ala Thr Val Glu Gly Leu Lys
 130 135 140

Gln Leu Lys Arg Ala Glu Glu Ile Ala Ala Leu Arg Gly Ile Ser Val
 145 150 155 160

Ser Asp Leu Ala

<210> 610

<211> 146

<212> PRT

<213> Streptococcus pneumoniae

<400> 610

Met Lys Leu His Glu Leu Lys Pro Ala Glu Gly Ser Arg Lys Val Arg
 1 5 10 15

Asn Arg Val Gly Arg Gly Thr Ser Ser Gly Asn Gly Lys Thr Ser Gly
 20 25 30

Arg Gly Gln Lys Gly Gln Lys Ala Arg Ser Gly Gly Gly Val Arg Leu
 35 40 45

Gly Phe Glu Gly Gly Gln Thr Pro Leu Phe Arg Arg Leu Pro Lys Arg
 50 55 60

Gly Phe Thr Asn Ile Asn Ala Lys Glu Tyr Ala Ile Val Asn Leu Asp
 65 70 75 80

Gln Leu Asn Val Phe Glu Asp Gly Ala Glu Val Thr Pro Val Val Leu
 85 90 95

Ile Glu Ala Gly Ile Val Lys Ala Glu Lys Ser Gly Ile Lys Ile Leu
 100 105 110

Gly Asn Gly Glu Leu Thr Lys Lys Leu Thr Val Lys Ala Ala Lys Phe
 115 120 125

Ser Lys Ser Ala Glu Glu Ala Ile Thr Ala Lys Gly Gly Ser Val Glu
 130 135 140

Val Ile
 145

<210> 611
 <211> 436
 <212> PRT
 <213> Streptococcus pneumoniae

<400> 611

Met Phe Phe Lys Leu Leu Arg Glu Ala Leu Lys Val Lys Gln Val Arg
 1 5 10 15

Ser Lys Ile Leu Phe Thr Ile Phe Ile Val Leu Val Phe Arg Ile Gly
 20 25 30

Thr Ser Ile Thr Val Pro Gly Val Asn Ala Asn Ser Leu Asn Ala Leu
 35 40 45

Ser Gly Leu Ser Phe Leu Asn Met Leu Ser Leu Val Ser Gly Asn Ala
 50 55 60

Leu Lys Asn Phe Ser Ile Phe Ala Leu Gly Val Ser Pro Tyr Ile Thr
 65 70 75 80

Ala Ser Ile Val Val Gln Leu Leu Gln Met Asp Ile Leu Pro Lys Phe
 85 90 95

Val Glu Trp Gly Lys Gln Gly Glu Val Gly Arg Arg Lys Leu Asn Gln
 100 105 110

Ala Thr Arg Tyr Ile Ala Leu Val Leu Ala Phe Val Gln Ser Ile Gly
 115 120 125

Ile Thr Ala Gly Phe Asn Thr Leu Ala Gly Ala Gln Leu Ile Lys Thr
 130 135 140

Ala Leu Thr Pro Gln Val Phe Leu Thr Ile Gly Ile Ile Leu Thr Ala
 145 150 155 160

Gly Ser Met Ile Val Thr Trp Leu Gly Glu Gln Ile Thr Asp Lys Gly
 165 170 175

Tyr Gly Asn Gly Val Ser Met Ile Ile Phe Ala Gly Ile Val Ser Ser
 180 185 190

Ile Pro Glu Met Ile Gln Gly Ile Tyr Val Asp Tyr Phe Val Asn Val
 195 200 205

Pro Ser Ser Arg Ile Thr Ser Ser Ile Ile Phe Val Ile Ile Leu Ile
 210 215 220

Ile Thr Val Leu Leu Ile Ile Tyr Phe Thr Thr Tyr Val Gln Gln Ala
 225 230 235 240

Glu Tyr Lys Ile Pro Ile Gln Tyr Thr Lys Val Ala Gln Gly Ala Pro
 245 250 255

Ser Ser Ser Tyr Leu Pro Leu Lys Val Asn Pro Ala Gly Val Ile Pro
 260 265 270

Val Ile Phe Ala Ser Ser Ile Thr Ala Ala Pro Ala Ala Ile Leu Gln
 275 280 285

Phe Leu Ser Ala Thr Gly His Asp Trp Ala Trp Val Arg Val Ala Gln
 290 295 300

Glu Met Leu Ala Thr Thr Ser Pro Thr Gly Ile Ala Met Tyr Ala Leu
 305 310 315 320

Leu Ile Ile Leu Phe Thr Phe Phe Tyr Thr Phe Val Gln Ile Asn Pro
 325 330 335

Glu Lys Ala Ala Glu Ser Leu Gln Lys Ser Gly Ala Tyr Ile His Gly
 340 345 350

Val Arg Pro Gly Lys Gly Thr Glu Glu Tyr Met Ser Lys Leu Leu Arg
 355 360 365

Arg Leu Ala Thr Val Gly Ser Leu Phe Leu Gly Val Ile Ser Ile Leu
 370 375 380

Pro Ile Ala Ala Lys Asp Val Phe Gly Leu Ser Asp Val Val Ala Phe
 385 390 395 400

Gly Gly Thr Ser Leu Leu Ile Ile Ile Ser Thr Gly Ile Glu Gly Ile
 405 410 415

Lys Gln Leu Glu Gly Tyr Leu Leu Lys Arg Lys Tyr Val Gly Phe Met
 420 425 430

Asp Arg Thr Glu
 435

<210> 612
 <211> 212
 <212> PRT
 <213> Streptococcus pneumoniae

<400> 612

Met Asn Leu Leu Ile Met Gly Leu Pro Gly Ala Gly Lys Gly Thr Gln
 1 5 10 15

Ala Ala Lys Ile Val Glu Gln Phe His Val Ala His Ile Ser Thr Gly
 20 25 30

Asp Met Phe Arg Ala Ala Met Ala Asn Gln Thr Glu Met Gly Val Leu
 35 40 45

Ala Lys Ser Tyr Ile Asp Lys Gly Glu Leu Val Pro Asp Glu Val Thr
 50 55 60

Asn Gly Ile Val Lys Glu Arg Leu Ser Gln Asp Asp Ile Lys Glu Thr
 65 70 75 80

Gly Phe Leu Leu Asp Gly Tyr Pro Arg Thr Ile Glu Gln Ala His Ala
 85 90 95

Leu Asp Lys Thr Leu Ala Glu Leu Gly Ile Glu Leu Glu Gly Val Ile
100 105 110

Asn Ile Glu Val Asn Pro Asp Ser Leu Leu Glu Arg Leu Ser Gly Arg
115 120 125

Ile Ile His Arg Val Thr Gly Glu Thr Phe His Lys Val Phe Asn Pro
130 135 140

Pro Val Asp Tyr Lys Glu Glu Asp Tyr Tyr Gln Arg Glu Asp Asp Lys
145 150 155 160

Pro Glu Thr Val Lys Arg Arg Leu Asp Val Asn Ile Ala Gln Gly Glu
165 170 175

Pro Ile Ile Ala His Tyr Arg Ala Lys Gly Leu Val His Asp Ile Glu
180 185 190

Gly Asn Gln Asp Ile Asn Asp Val Phe Ser Asp Ile Glu Lys Val Leu
195 200 205

Thr Asn Leu Lys
210

<210> 613

<211> 121

<212> FRT

<213> Streptococcus pneumoniae

<400> 613

Met Ala Arg Ile Ala Gly Val Asp Ile Pro Asn Asp Lys Arg Val Val
1 5 10 15

Ile Ser Leu Thr Tyr Val Tyr Gly Ile Gly Leu Ala Thr Ser Lys Lys
20 25 30

Ile Leu Ala Ala Ala Gly Ile Ser Glu Asp Val Arg Val Arg Asp Leu
35 40 45

Thr Ser Asp Gln Glu Asp Ala Ile Arg Arg Glu Val Asp Ala Ile Lys
50 55 60

Val Glu Gly Asp Leu Arg Arg Glu Val Asn Leu Asn Ile Lys Arg Leu
65 70 75 80

Met Glu Ile Gly Ser Tyr Arg Gly Ile Arg His Arg Arg Gly Leu Pro
85 90 95

Val Arg Gly Gln Asn Thr Lys Asn Asn Ala Arg Thr Arg Lys Gly Lys
100 105 110

Ala Val Ala Ile Ala Gly Lys Lys Lys
115 120

<210> 614
<211> 127
<212> PRT
<213> Streptococcus pneumoniae

<400> 614

Met Ala Lys Pro Thr Arg Lys Arg Arg Val Lys Lys Asn Ile Glu Ser
1 5 10 15

Gly Ile Ala His Ile His Ala Thr Phe Asn Asn Thr Ile Val Met Ile
20 25 30

Thr Asp Val His Gly Asn Ala Ile Ala Trp Ser Ser Ala Gly Ala Leu
35 40 45

Gly Phe Lys Gly Ser Arg Lys Ser Thr Pro Phe Ala Ala Gln Met Ala
50 55 60

Ser Glu Ala Ala Ala Lys Ser Ala Gln Glu His Gly Leu Lys Ser Val
65 70 75 80

Glu Val Thr Val Lys Gly Pro Gly Ser Gly Arg Glu Ser Ala Ile Arg
85 90 95

Ala Leu Ala Ala Ala Gly Leu Glu Val Thr Ala Ile Arg Asp Val Thr
100 105 110

Pro Val Pro His Asn Gly Ala Arg Pro Pro Lys Arg Arg Arg Val
115 120 125

<210> 615
 <211> 128
 <212> PRT
 <213> Streptococcus pneumoniae
 <400> 615
 Met Ala Tyr Arg Lys Leu Gly Arg Thr Ser Ser Gln Arg Lys Ala Met
 1 5 10 15
 Leu Arg Asp Leu Thr Thr Asp Leu Leu Ile Asn Glu Ser Ile Val Thr
 20 25 30
 Thr Glu Ala Arg Ala Lys Glu Ile Arg Lys Thr Val Glu Lys Met Ile
 35 40 45
 Thr Leu Gly Lys Arg Gly Asp Leu His Ala Arg Arg Gln Ala Ala Ala
 50 55 60
 Phe Val Arg Asn Glu Ile Ala Ser Glu Asn Tyr Asp Glu Ala Thr Asp
 65 70 75 80
 Lys Tyr Thr Ser Thr Thr Ala Leu Gln Lys Leu Phe Ser Glu Ile Ala
 85 90 95
 Pro Arg Tyr Ala Glu Arg Asn Gly Gly Tyr Thr Arg Ile Leu Lys Thr
 100 105 110
 Glu Ser Arg Arg Gly Asp Ala Ala Pro Met Ala Ile Ile Glu Leu Val
 115 120 125
 <210> 616
 <211> 137
 <212> PRT
 <213> Streptococcus pneumoniae
 <400> 616
 Met Pro Thr Ile Asn Gln Leu Val Arg Lys Pro Arg Lys Ser Lys Val
 1 5 10 15
 Glu Lys Ser Lys Ser Pro Ala Leu Asn Val Gly Tyr Asn Ser His Lys
 20 25 30

Lys Val Gln Thr Asn Val Ser Ser Pro Gln Lys Arg Gly Val Ala Thr
 35 40 45

Arg Val Gly Thr Met Thr Pro Lys Lys Pro Asn Ser Ala Leu Arg Lys
 50 55 60

Phe Ala Arg Val Arg Leu Ser Asn Leu Ile Glu Val Thr Ala Tyr Ile
 65 70 75 80

Pro Gly Ile Gly His Asn Leu Gln Glu His Ser Val Val Leu Leu Arg
 85 90 95

Gly Gly Arg Val Lys Asp Leu Pro Gly Val Arg Tyr His Ile Val Arg
 100 105 110

Gly Ala Leu Asp Thr Ala Gly Val Asn Asp Arg Lys Gln Gly Arg Ser
 115 120 125

Lys Tyr Gly Thr Lys Arg Pro Lys Ala
 130 135

<210> 617

<211> 156

<212> PRT

<213> Streptococcus pneumoniae

<400> 617

Met Ser Arg Lys Asn Arg Ala Pro Lys Arg Asp Val Leu Pro Asp Pro
 1 5 10 15

Leu Tyr Asn Ser Gln Leu Val Thr Arg Leu Ile Asn Arg Val Met Leu
 20 25 30

Asp Gly Lys Arg Gly Thr Ala Ala Ser Ile Val Tyr Gly Ala Phe Glu
 35 40 45

Gln Ile Lys Glu Ala Thr Gly Asn Asp Ala Leu Glu Val Phe Glu Thr
 50 55 60

Ala Met Glu Asn Ile Met Pro Val Leu Glu Val Arg Ala Arg Arg Val
 65 70 75 80

Gly Gly Ser Asn Tyr Gln Val Pro Val Lys Val Arg Pro Glu Arg Arg
85 90 95

Thr Thr Leu Gly Leu Arg Trp Leu Val Thr Ile Ala Arg Leu Arg Gly
100 105 110

Glu His Thr Met Gln Asp Arg Leu Ala Lys Glu Ile Leu Asp Ala Ala
115 120 125

Asn Asn Thr Gly Ala Ala Val Lys Lys Arg Glu Asp Thr His Arg Met
130 135 140

Ala Glu Ala Asn Arg Ala Phe Ala His Phe Arg Trp
145 150 155

<210> 618

<211> 303

<212> PRT

<213> Streptococcus pneumoniae

<400> 618

Met Thr Glu Lys Leu Gln Leu Thr Lys Ser Asp Arg Lys Lys Val Trp
1 5 10 15

Trp Arg Ser Thr Phe Leu Gln Gly Ser Trp Asn Phe Glu Arg Met Gln
20 25 30

Asn Leu Gly Trp Ala Tyr Thr Leu Ile Pro Ala Ile Lys Lys Leu Tyr
35 40 45

Thr Lys Lys Glu Asp Gln Ile Ala Ala Leu Glu Arg His Leu Glu Phe
50 55 60

Phe Asn Thr His Pro Tyr Val Ala Ala Pro Val Met Gly Val Thr Leu
65 70 75 80

Ala Leu Glu Glu Glu Arg Ala Asn Gly Val Glu Ile Asp Asp Ala Ala
85 90 95

Ile Gln Gly Val Lys Ile Gly Met Met Gly Pro Leu Ala Gly Ile Gly
100 105 110

Asp Pro Val Phe Trp Phe Thr Val Arg Pro Ile Leu Gly Ser Leu Gly
 115 120 125
 Ala Ser Leu Ala Leu Thr Gly Asn Ile Leu Gly Pro Leu Leu Phe Phe
 130 135 140
 Val Ala Trp Asn Leu Ile Arg Met Ser Phe Leu Trp Tyr Val Gln Glu
 145 150 155 160
 Ile Gly Tyr Lys Ala Gly Ser Glu Ile Thr Lys Asp Met Ser Gly Gly
 165 170 175
 Ile Leu Gln Asp Ile Thr Lys Gly Ala Ser Ile Leu Gly Met Phe Ile
 180 185 190
 Leu Ala Val Leu Val Gln Arg Trp Val Asn Ile Lys Phe Ala Phe Asp
 195 200 205
 Val Ser Lys Val Gln Leu Asp Glu Lys Ala Tyr Ile His Trp Asp Lys
 210 215 220
 Leu Pro Glu Gly Ser Lys Gly Ile Gln Glu Ala Phe Ala Gln Val Gly
 225 230 235 240
 Gln Gly Leu Ser Gln Thr Pro Glu Lys Val Thr Thr Phe Gln Gln Asn
 245 250 255
 Leu Asp Met Leu Ile Pro Gly Leu Ser Gly Leu Leu Leu Thr Leu Leu
 260 265 270
 Cys Met Tyr Leu Leu Lys Lys Lys Val Ser Pro Ile Thr Ile Ile Leu
 275 280 285
 Ala Leu Phe Ala Val Gly Ile Val Ala His Val Leu His Ile Met
 290 295 300

<210> 619
 <211> 267
 <212> PRT
 <213> Streptococcus pneumoniae
 <400> 619

Met Ser Ile Ile Ser Met Val Leu Val Val Val Val Ala Phe Phe Ala
 1 5 10 15
 Gly Leu Glu Gly Ile Leu Asp Gln Phe Gln Phe His Gln Pro Leu Val
 20 25 30
 Ala Cys Thr Leu Ile Gly Leu Val Thr Gly His Leu Glu Ala Gly Ile
 35 40 45
 Ile Leu Gly Gly Ser Leu Gln Met Ile Ala Leu Gly Trp Ser Asn Ile
 50 55 60
 Gly Ala Ala Ile Ala Pro Asp Ala Ala Leu Ala Ser Val Ala Ala Ala
 65 70 75 80
 Ile Ile Met Val Leu Gly Gly Asp Phe Thr Lys Thr Gly Ile Gly Val
 85 90 95
 Ala Gln Ala Val Ala Ile Pro Leu Ala Val Ala Gly Leu Phe Leu Thr
 100 105 110
 Met Ile Val Arg Thr Ile Ser Val Gly Leu Val His Thr Ala Asp Ala
 115 120 125
 Ala Ala Lys Lys Gly Asp Phe Gly Ala Val Glu Arg Ala His Phe Ile
 130 135 140
 Ala Leu Leu Phe Gln Gly Leu Arg Ile Ala Leu Pro Ala Ala Leu Leu
 145 150 155 160
 Leu Met Val Pro Thr Glu Thr Val Gln Ser Ile Leu Ser Ala Met Pro
 165 170 175
 Asp Trp Leu Lys Asp Gly Met Ala Ile Gly Gly Gly Met Val Val Ala
 180 185 190
 Val Gly Tyr Ala Met Val Ile Asn Met Met Ala Thr Arg Glu Val Trp
 195 200 205
 Pro Phe Phe Ala Leu Gly Phe Val Leu Ala Ala Val Ser Asp Ile Thr
 210 215 220

Leu Ile Gly Phe Gly Ala Ile Gly Val Ala Ile Ala Leu Ile Tyr Leu
 225 230 235 240

His Leu Ser Lys Thr Gly Gly Asn Gly Gly Gly Gly Ala Ala Thr Ser
 245 250 255

Asn Asp Pro Ile Gly Asp Ile Leu Glu Asp Tyr
 260 265

<210> 620

<211> 332

<212> PRT

<213> Streptococcus pneumoniae

<400> 620

Met Thr Ile Met Ser Ile Gly Ile Ile Ile Ala Ser His Gly Glu Phe
 1 5 10 15

Ala Ala Gly Ile His Gln Ser Gly Ser Met Ile Phe Gly Glu Gln Glu
 20 25 30

Lys Val Gln Val Val Thr Phe Met Pro Asn Glu Gly Pro Asp Asp Leu
 35 40 45

Tyr Ala Lys Phe Asn Asn Ala Val Ala Ala Phe Asp Ala Glu Asp Glu
 50 55 60

Val Leu Val Leu Ala Asp Leu Trp Ser Gly Ser Pro Phe Asn Gln Ala
 65 70 75 80

Ser Arg Val Met Gly Glu Asn Pro Glu Arg Lys Phe Ala Ile Ile Thr
 85 90 95

Gly Leu Asn Leu Pro Met Leu Ile Gln Ala Tyr Thr Glu Arg Leu Met
 100 105 110

Asp Ala Ala Ala Gly Val Glu Lys Val Ala Ala Asn Ile Ile Lys Glu
 115 120 125

Ala Lys Asp Gly Ile Lys Ala Leu Pro Glu Glu Leu Asn Pro Val Glu
 130 135 140

Glu Val Ala Ser Ala Ala Ala Ala Pro Val Ala Gln Thr Ala Ile Pro
 145 150 155 160

Glu Gly Thr Val Ile Gly Asp Gly Lys Leu Lys Ile Asn Leu Ala Arg
 165 170 175

Leu Asp Thr Arg Leu Leu His Gly Gln Val Ala Thr Ala Trp Thr Pro
 180 185 190

Asp Ser Lys Ala Asn Arg Ile Ile Val Ala Ser Asp Asn Val Ala Lys
 195 200 205

Asp Asp Leu Arg Lys Glu Leu Ile Lys Gln Ala Ala Pro Gly Asn Val
 210 215 220

Lys Ala Asn Val Val Pro Ile Gln Lys Leu Ile Glu Ile Ser Lys Asp
 225 230 235 240

Pro Arg Phe Gly Glu Thr His Ala Leu Ile Leu Phe Glu Thr Pro Gln
 245 250 255

Asp Ala Leu Arg Ala Ile Glu Gly Gly Val Pro Ile Lys Thr Leu Asn
 260 265 270

Val Gly Ser Met Ala His Ser Thr Gly Lys Thr Leu Val Asn Thr Val
 275 280 285

Leu Ser Met Asp Lys Glu Asp Val Ala Thr Phe Glu Lys Met Arg Asp
 290 295 300

Leu Gly Val Glu Phe Asp Val Arg Lys Val Pro Asn Asp Ser Lys Lys
 305 310 315 320

Asp Leu Phe Asp Leu Ile Asn Lys Ala Asn Val Lys
 325 330

<210> 621

<211> 339

<212> PRT

<213> Streptococcus pneumoniae

<400> 621

Met Lys Ala Val Val Val Asn Pro Glu Ser Thr Gly Val Ala Ile Glu
 1 5 10 15

Glu Lys Val Leu Arg Pro Leu Glu Thr Gly Glu Ala Leu Val Glu Val
 20 25 30

Glu Tyr Cys Gly Val Cys His Thr Asp Leu His Val Ala His Gly Asp
 35 40 45

Phe Gly Gln Val Pro Gly Arg Val Leu Gly His Glu Gly Ile Gly Ile
 50 55 60

Val Lys Glu Ile Ala Pro Asp Val Lys Ser Leu Lys Val Gly Asp Arg
 65 70 75 80

Val Ser Val Ala Trp Phe Phe Glu Gly Cys Gly Thr Cys Glu Tyr Cys
 85 90 95

Thr Thr Gly Arg Glu Thr Leu Cys Arg Thr Val Lys Asn Ala Gly Tyr
 100 105 110

Ser Val Asp Gly Gly Met Ala Glu Gln Cys Ile Val Thr Ala Asp Tyr
 115 120 125

Ala Val Lys Val Pro Asp Gly Leu Asp Pro Ala Gln Ala Ser Ser Ile
 130 135 140

Thr Cys Ala Gly Val Thr Thr Tyr Lys Ala Ile Lys Glu Ala Lys Val
 145 150 155 160

Glu Pro Gly Gln Trp Val Val Leu Tyr Gly Ala Gly Gly Leu Gly Asn
 165 170 175

Leu Ala Val Gln Tyr Ala Lys Lys Val Phe Asn Ala His Val Ile Ala
 180 185 190

Val Asp Ile Asn Asn Asp Lys Leu Ala Leu Ala Lys Glu Val Gly Ala
 195 200 205

Asp Ile Val Ile Asn Gly Leu Glu Val Glu Asp Val Ala Gly Leu Ile
 210 215 220

Lys Glu Lys Thr Asp Gly Gly Ala His Ser Ala Val Val Thr Ala Val
 225 230 235 240

Ser Lys Val Ala Phe Asn Gln Ala Val Asp Ser Ile Arg Ala Gly Gly
 245 250 255

Arg Val Val Ala Val Gly Leu Pro Ser Glu Met Met Glu Leu Ser Ile
 260 265 270

Val Lys Thr Val Leu Asp Gly Ile Gln Val Ile Gly Ser Leu Val Gly
 275 280 285

Thr Arg Lys Asp Leu Glu Glu Ala Phe Gln Phe Gly Ala Glu Gly Leu
 290 295 300

Val Val Pro Val Val Gln Lys Arg Pro Val Glu Asp Ala Val Ala Ile
 305 310 315 320

Phe Asp Glu Met Glu Lys Gly Gln Ile Gln Gly Arg Met Val Leu Asp
 325 330 335

Phe Thr His

<210> 622
 <211> 148
 <212> PRT
 <213> Streptococcus pneumoniae

<400> 622

Met Asn Lys Thr Thr Phe Met Ala Lys Pro Gly Gln Val Glu Arg Lys
 1 5 10 15

Trp Tyr Val Val Asp Ala Thr Asp Val Pro Leu Gly Arg Leu Ser Ala
 20 25 30

Val Val Ala Ser Val Leu Arg Gly Lys Asn Lys Pro Thr Phe Thr Pro
 35 40 45

His Thr Asp Thr Gly Asp Phe Val Ile Val Ile Asn Ala Glu Lys Val
 50 55 60

Lys Leu Thr Gly Lys Lys Ala Thr Asp Lys Ile Tyr Tyr Thr His Ser
65 70 75 80

Asn His Pro Gly Gly Leu Lys Gln Ile Ser Ala Gly Glu Leu Arg Ser
85 90 95

Lys Asn Ala Val Arg Leu Ile Glu Lys Ser Val Lys Gly Met Leu Pro
100 105 110

His Asn Thr Leu Gly Arg Ala Gln Gly Met Lys Leu Lys Val Phe Val
115 120 125

Gly Ala Glu His Thr His Ala Ala Gln Gln Pro Glu Val Leu Asp Ile
130 135 140

Ser Gly Leu Ile
145

<210> 623
<211> 130
<212> PRT
<213> Streptococcus pneumoniae

<400> 623

Met Ser Gln Ala Gln Tyr Ala Gly Thr Gly Arg Arg Lys Asn Ala Val
1 5 10 15

Ala Arg Val Arg Leu Val Pro Gly Thr Gly Lys Ile Thr Val Asn Lys
20 25 30

Lys Asp Val Glu Glu Tyr Ile Pro His Ala Asp Leu Arg Leu Val Ile
35 40 45

Asn Gln Pro Phe Ala Val Thr Ser Thr Val Gly Ser Tyr Asp Val Phe
50 55 60

Val Asn Val Ile Gly Gly Gly Tyr Ala Gly Gln Ser Gly Ala Ile Arg
65 70 75 80

His Gly Ile Ala Arg Ala Leu Leu Gln Val Asp Pro Asp Phe Arg Asp
85 90 95

Ser Leu Lys Arg Ala Gly Leu Leu Thr Arg Asp Ser Arg Lys Val Glu
 100 105 110

Arg Lys Lys Pro Gly Leu Lys Lys Ala Arg Lys Ala Ser Gln Phe Ser
 115 120 125

Lys Arg
 130

<210> 624
 <211> 272
 <212> PRT
 <213> Streptococcus pneumoniae

<400> 624

Met Thr Gly Ser Asn Lys Leu Thr Lys Arg Asp Tyr Leu Lys Thr Ser
 1 5 10 15

Leu Arg Ala Phe Phe Cys Gln Asn Gly Phe Asn Tyr Ser Asn Tyr Gln
 20 25 30

Gly Leu Gly Tyr Ala Asn Val Met Tyr Pro Ala Leu Lys Lys His Tyr
 35 40 45

Gly Glu Asp Gln Glu Gly Phe Tyr Gln Ala Leu Glu Glu Asn Cys Glu
 50 55 60

Phe Tyr Asn Thr Asn Pro His Phe Leu Pro Phe Ile Thr Ser Leu His
 65 70 75 80

Leu Val Met Leu Glu Asn Gly Arg Pro Ala Lys Glu Thr Arg Ser Ile
 85 90 95

Lys Met Ala Leu Met Gly Pro Leu Ala Gly Ile Gly Asp Ser Leu Ser
 100 105 110

Gln Phe Cys Leu Ala Pro Leu Phe Ser Thr Ile Ala Ala Ser Phe Ala
 115 120 125

Gln Glu Gly Leu Val Val Gly Pro Ile Leu Phe Phe Leu Ala Met Asn
 130 135 140

Thr Ile Leu Thr Ala Ile Lys Leu Ser Thr Gly Leu Tyr Gly Tyr Lys
145 150 155 160

Leu Gly Thr Thr Val Ile Asp Lys Leu Ser Glu Gln Met Ala Thr Ile
165 170 175

Ser Arg Ile Ala Asn Ile Ile Gly Val Thr Val Ile Ala Gly Leu Ala
180 185 190

Ala Thr Ser Val Lys Ile Met Val Pro Ile Thr Phe Ala Ala Gly Glu
195 200 205

Val Lys Ala Asp Ala Lys Gln Ser Ile Val Ser Ile Gln Gly Met Leu
210 215 220

Asp Lys Val Ala Pro Ala Leu Leu Pro Ala Leu Phe Thr Leu Leu Val
225 230 235 240

Tyr Tyr Leu Ile Lys Glu Lys Lys Trp Thr Thr Tyr Lys Leu Val Ile
245 250 255

Leu Thr Val Ile Ile Gly Ile Ile Gly Ser Trp Leu Lys Ile Ile Ala
260 265 270

<210> 625

<211> 494

<212> PRT

<213> Streptococcus pneumoniae

<400> 625

Met Lys Lys Gln Ala Phe Ser Ser Glu Gln Tyr Leu Asn Leu Gln Arg
1 5 10 15

Asp His Ile Leu Glu Arg Ile Asn Gln Phe Asp Gly Lys Leu Tyr Leu
20 25 30

Glu Phe Gly Gly Lys Met Leu Glu Asp Phe His Ala Ala Arg Val Leu
35 40 45

Pro Gly Tyr Glu Pro Asp Asn Lys Ile Lys Leu Leu Gln Glu Leu Lys
50 55 60

Glu Gln Val Glu Val Val Ile Ala Ile Asn Ala Ser Asn Ile Glu His
 65 70 75 80

Ser Lys Ala Arg Gly Asp Leu Gly Ile Ser Tyr Asp Gln Glu Val Leu
 85 90 95

Arg Leu Ile Asp Lys Phe Asn Glu Leu Gly Ile Phe Val Gly Ser Val
 100 105 110

Val Ile Thr Gln Tyr Ala Gly Gln Pro Ala Ala Asp Ala Phe Arg Asn
 115 120 125

Gln Leu Glu Lys Asn Gly Ile Asp Ser Tyr Leu His Tyr Pro Ile Lys
 130 135 140

Gly Tyr Pro Thr Asp Met Asp His Ile Ile Ser Pro Glu Gly Met Gly
 145 150 155 160

Lys Asn Asp Tyr Ile Lys Thr Ser Arg Asn Leu Ile Val Val Thr Ala
 165 170 175

Pro Gly Pro Gly Ser Gly Lys Leu Ala Thr Cys Met Ser Asn Met Tyr
 180 185 190

His Asp Gln Ile Asn Gly Ile Lys Ser Gly Tyr Ala Lys Phe Glu Thr
 195 200 205

Phe Pro Val Trp Asn Leu Pro Leu His His Pro Val Asn Leu Ala Tyr
 210 215 220

Glu Ala Ala Thr Ala Asp Leu Asp Asp Val Asn Met Ile Asp Pro Phe
 225 230 235 240

His Leu Gln Thr Tyr Gly Glu Thr Thr Val Asn Tyr Asn Arg Asp Ile
 245 250 255

Glu Ile Phe Pro Val Leu Lys Arg Met Leu Glu Arg Ile Leu Gly Lys
 260 265 270

Ser Pro Tyr Ala Ser Pro Thr Asp Met Gly Val Asn Met Val Gly Phe

275	280	285
Ala Ile Thr Asp Asp Glu 290	Ala Ala Val Glu Ala 295	Ser Lys Gln Glu Ile 300
Ile Arg Arg Tyr Tyr Gln 305	Thr Val Leu Asp 310	Phe Lys Ala Glu Lys Val 315 320
Gly Glu Ala Ala Val Lys Lys 325	Ile Glu Leu Leu Met 330	Asn Asp Leu Gly 335
Ile Thr Pro Ala Asp Arg Lys 340	Val Ala Val Val Ala 345	Arg Gln Lys Ala 350
Glu Glu Thr Gly Gly Pro Ala 355	Leu Ala Phe Glu Leu 360	Pro Asn Gly Glu 365
Ile Ile Thr Gly Lys Asn Ser 370	Glu Leu Phe Gly 375	Pro Thr Ala Ala Ala 380
Leu Ile Asn Ala Ile Lys Lys 385	Ser Ala Asp 390	Ile Ala Lys Glu Val Lys 395 400
Leu Ile Glu Pro Glu Val Val 405	Lys Pro Ile Gln Gly Leu Lys 410	Ile Asp 415
His Leu Gly Ser Arg Asn Pro 420	Arg Leu His Ser Asn Glu 425	Ile Leu Ile 430
Ala Leu Ala Ile Thr Ala Thr 435	Glu Asn Pro Asp Ala Ala 440	Arg Ala Met 445
Glu Glu Leu Gly Asn Leu Lys 450	Gly Ser Glu Ala His 455	Ser Thr Ile Ile 460
Leu Thr Asp Glu Asp Lys Asn 465	Val Leu Arg Lys Leu Gly 470	Ile Asn Val 475 480
Thr Phe Asp Pro Tyr Tyr Gln 485	Tyr Asp Arg Leu Tyr Arg 490	Lys

<210> 626
 <211> 719
 <212> PRT
 <213> Streptococcus pneumoniae

 <400> 626

 Met Asn Lys Pro Thr Ile Leu Arg Leu Ile Lys Tyr Leu Ser Ile Ser
 1 5 10 15

 Phe Leu Ser Leu Val Ile Ala Ala Ile Val Leu Gly Gly Gly Val Phe
 20 25 30

 Phe Tyr Tyr Val Ser Lys Ala Pro Ser Leu Ser Glu Ser Lys Leu Val
 35 40 45

 Ala Thr Thr Ser Ser Lys Ile Tyr Asp Asn Lys Asn Gln Leu Ile Ala
 50 55 60

 Asp Leu Gly Ser Glu Arg Arg Val Asn Ala Gln Ala Asn Asp Ile Pro
 65 70 75 80

 Thr Asp Leu Val Lys Ala Ile Val Ser Ile Glu Asp His Arg Phe Phe
 85 90 95

 Asp His Arg Gly Ile Asp Thr Ile Arg Ile Leu Gly Ala Phe Leu Arg
 100 105 110

 Asn Leu Gln Ser Asn Ser Leu Gln Gly Gly Ser Thr Leu Thr Gln Gln
 115 120 125

 Leu Ile Lys Leu Thr Tyr Phe Ser Thr Ser Thr Ser Asp Gln Thr Ile
 130 135 140

 Ser Arg Lys Ala Gln Glu Ala Trp Leu Ala Ile Gln Leu Glu Gln Lys
 145 150 155 160

 Ala Thr Lys Gln Glu Ile Leu Thr Tyr Tyr Ile Asn Lys Val Tyr Met
 165 170 175

 Ser Asn Gly Asn Tyr Gly Met Gln Thr Ala Ala Gln Asn Tyr Tyr Gly
 180 185 190

Lys Asp Leu Asn Asn Leu Ser Leu Pro Gln Leu Ala Leu Leu Ala Gly
 195 200 205
 Met Pro Gln Ala Pro Asn Gln Tyr Asp Pro Tyr Ser His Pro Glu Ala
 210 215 220
 Ala Gln Asp Arg Arg Asn Leu Val Leu Ser Glu Met Lys Asn Gln Gly
 225 230 235 240
 Tyr Ile Ser Ala Glu Gln Tyr Glu Lys Ala Val Asn Thr Pro Ile Thr
 245 250 255
 Asp Gly Leu Gln Ser Leu Lys Ser Ala Ser Asn Tyr Pro Ala Tyr Met
 260 265 270
 Asp Asn Tyr Leu Lys Glu Val Ile Asn Gln Val Glu Glu Thr Gly
 275 280 285
 Tyr Asn Leu Leu Thr Thr Gly Met Asp Val Tyr Thr Asn Val Asp Gln
 290 295 300
 Glu Ala Gln Lys His Leu Trp Asp Ile Tyr Asn Thr Asp Glu Tyr Val
 305 310 315 320
 Ala Tyr Pro Asp Asp Glu Leu Gln Val Ala Ser Thr Ile Val Asp Val
 325 330 335
 Ser Asn Gly Lys Val Ile Ala Gln Leu Gly Ala Arg His Gln Ser Ser
 340 345 350
 Asn Val Ser Phe Gly Ile Asn Gln Ala Val Glu Thr Asn Arg Asp Trp
 355 360 365
 Gly Ser Thr Met Lys Pro Ile Thr Asp Tyr Ala Pro Ala Leu Glu Tyr
 370 375 380
 Gly Val Tyr Asp Ser Thr Ala Thr Ile Val His Asp Glu Pro Tyr Asn
 385 390 395 400
 Tyr Pro Gly Thr Asn Thr Pro Val Tyr Asn Trp Asp Arg Gly Tyr Phe
 405 410 415

Gly Asn Ile Thr Leu Gln Tyr Ala Leu Gln Gln Ser Arg Asn Val Pro
 420 425 430
 Ala Val Glu Thr Leu Asn Lys Val Gly Leu Asn Arg Ala Lys Thr Phe
 435 440 445
 Leu Asn Gly Leu Gly Ile Asp Tyr Pro Ser Ile His Tyr Ser Asn Ala
 450 455 460
 Ile Ser Ser Asn Thr Thr Glu Ser Asp Lys Lys Tyr Gly Ala Ser Ser
 465 470 475 480
 Glu Lys Met Ala Ala Ala Tyr Ala Ala Phe Ala Asn Gly Gly Thr Tyr
 485 490 495
 Tyr Lys Pro Met Tyr Ile His Lys Val Val Phe Ser Asp Gly Ser Glu
 500 505 510
 Lys Glu Phe Ser Asn Val Gly Thr Arg Ala Met Lys Glu Thr Thr Ala
 515 520 525
 Tyr Met Met Thr Asp Met Met Lys Thr Val Leu Thr Tyr Gly Thr Gly
 530 535 540
 Arg Asn Ala Tyr Leu Ala Trp Leu Pro Gln Ala Gly Lys Thr Gly Thr
 545 550 555 560
 Ser Asn Tyr Thr Asp Glu Glu Ile Glu Asn His Ile Lys Thr Ser Gln
 565 570 575
 Phe Val Ala Pro Asp Glu Leu Phe Ala Gly Tyr Thr Arg Lys Tyr Ser
 580 585 590
 Met Ala Val Trp Thr Gly Tyr Ser Asn Arg Leu Thr Pro Leu Val Gly
 595 600 605
 Asn Gly Leu Thr Val Ala Ala Lys Val Tyr Arg Ser Met Met Thr Tyr
 610 615 620
 Leu Ser Glu Gly Ser Asn Pro Glu Asp Trp Asn Ile Pro Glu Gly Leu
 625 630 635 640

Tyr Arg Asn Gly Glu Phe Val Phe Lys Asn Gly Ala Arg Ser Thr Trp
645 650 655

Asn Ser Pro Ala Pro Gln Gln Pro Pro Ser Thr Glu Ser Ser Ser Ser
660 665 670

Ser Ser Asp Ser Ser Thr Ser Gln Ser Ser Ser Thr Thr Pro Ser Thr
675 680 685

Asn Asn Ser Thr Thr Thr Asn Pro Asn Asn Asn Thr Gln Gln Ser Asn
690 695 700

Thr Thr Pro Asp Gln Gln Asn Gln Asn Pro Gln Pro Ala Gln Pro
705 710 715

<210> 627

<211> 113

<212> PRT

<213> Streptococcus pneumoniae

<400> 627

Met Glu Arg Asp Met Ala Ser Ile Ile Phe Ser Ala Lys Asp Ile Phe
1 5 10 15

Glu Gln Glu Phe Gly Arg Glu Val Arg Gly Tyr Asn Lys Val Glu Val
20 25 30

Asp Glu Phe Leu Asp Asp Val Ile Lys Asp Tyr Glu Thr Tyr Ala Ala
35 40 45

Leu Val Lys Ser Leu Arg Gln Glu Ile Ala Asp Leu Lys Glu Glu Leu
50 55 60

Thr Arg Lys Pro Lys Pro Ser Pro Val Gln Ala Glu Pro Leu Glu Ala
65 70 75 80

Ala Ile Thr Ser Ser Met Thr Asn Phe Asp Ile Leu Lys Arg Leu Asn
85 90 95

Arg Leu Glu Lys Glu Val Phe Gly Lys Gln Ile Leu Asp Asn Ser Asp
100 105 110

Phe

<210> 628

<211> 464

<212> PRT

<213> Streptococcus pneumoniae

<400> 628

Met Ser Lys Lys Arg Arg Asn Arg His Lys Lys Glu Gly Gln Glu Pro
 1 5 10 15

Gln Phe Asp Phe Asp Glu Ala Lys Glu Leu Thr Val Gly Gln Ala Ile
 20 25 30

Arg Lys Asn Glu Glu Val Glu Ser Gly Val Leu Pro Glu Asp Ser Ile
 35 40 45

Leu Asp Lys Tyr Val Lys Gln His Arg Asp Glu Ile Glu Ala Asp Lys
 50 55 60

Phe Ala Thr Arg Gln Tyr Lys Lys Glu Glu Phe Val Glu Thr Gln Ser
 65 70 75 80

Leu Asp Asp Leu Ile Gln Glu Met Arg Glu Ala Val Glu Lys Ser Glu
 85 90 95

Ala Ser Ser Glu Glu Val Pro Ser Ser Glu Asp Ile Leu Leu Pro Leu
 100 105 110

Pro Leu Asp Asp Glu Glu Gln Gly Leu Asp Pro Leu Leu Leu Asp Asp
 115 120 125

Glu Asn Pro Thr Glu Met Thr Glu Glu Val Glu Glu Glu Gln Asn Leu
 130 135 140

Ser Arg Leu Asp Gln Glu Asp Ser Glu Lys Lys Ser Lys Lys Gly Phe
 145 150 155 160

Ile Leu Thr Val Leu Ala Leu Val Ser Val Ile Ile Cys Val Ser Ala
 165 170 175

Tyr Tyr Val Tyr Arg Gln Val Ala Arg Ser Thr Lys Glu Ile Glu Thr
 180 185 190
 Ser Gln Ser Thr Thr Ala Asn Gln Ser Asp Val Asp Asp Phe Asn Thr
 195 200 205
 Leu Tyr Asp Ala Phe Tyr Thr Asp Ser Asn Lys Thr Ala Leu Lys Asn
 210 215 220
 Ser Gln Phe Asp Lys Leu Ser Gln Leu Lys Thr Leu Leu Asp Lys Leu
 225 230 235 240
 Glu Gly Ser Arg Glu His Thr Leu Ala Lys Ser Lys Tyr Asp Ser Leu
 245 250 255
 Ala Thr Gln Ile Lys Ala Ile Gln Asp Val Asn Ala Gln Phe Glu Lys
 260 265 270
 Pro Ala Ile Val Asp Gly Val Leu Asp Thr Asn Ala Lys Ala Lys Ser
 275 280 285
 Asp Ala Lys Phe Thr Asp Ile Lys Thr Gly Asn Thr Glu Leu Asp Lys
 290 295 300
 Val Leu Asp Lys Ala Ile Ser Leu Gly Lys Ser Gln Gln Thr Ser Thr
 305 310 315 320
 Ser Ser Ser Ser Ser Ser Gln Thr Ser Ser Ser Ser Ser Ser Gln Ala
 325 330 335
 Ser Ser Asn Thr Thr Ser Glu Pro Lys Pro Ser Ser Ser Asn Glu Thr
 340 345 350
 Arg Ser Ser Arg Ser Glu Val Asn Met Gly Leu Ser Ser Ala Gly Val
 355 360 365
 Ala Val Gln Arg Ser Ala Ser Arg Val Ala Tyr Asn Gln Ser Ala Ile
 370 375 380
 Asp Asp Ser Asn Asn Ser Ala Trp Asp Phe Ala Asp Gly Val Leu Glu

385 390 395 400
 Gln Ile Leu Ala Thr Ser Arg Ser Arg Gly Tyr Ile Thr Gly Asp Gln
 405 410 415
 Tyr Ile Leu Glu Arg Val Asn Ile Val Asn Gly Asn Gly Tyr Tyr Asn
 420 425 430
 Leu Tyr Lys Pro Asp Gly Thr Tyr Leu Phe Thr Leu Asn Cys Lys Thr
 435 440 445
 Gly Tyr Phe Val Gly Asn Gly Ala Gly His Ala Asp Asp Leu Asp Tyr
 450 455 460

 <210> 629
 <211> 481
 <212> FRT
 <213> Streptococcus pneumoniae

 <400> 629
 Met Lys Gln Glu Glu Cys Gln Met Thr Lys Ala Asn Phe Gly Val Val
 1 5 10 15

 Gly Met Ala Val Met Gly Arg Asn Leu Ala Leu Asn Ile Glu Ser Arg
 20 25 30

 Gly Tyr Thr Val Ala Ile Tyr Asn Arg Ser Lys Glu Lys Thr Glu Asp
 35 40 45

 Val Ile Ala Cys His Pro Glu Lys Asn Phe Val Pro Ser Tyr Asp Val
 50 55 60

 Glu Ser Phe Val Asn Ser Ile Glu Lys Pro Arg Arg Ile Met Leu Met
 65 70 75 80

 Val Gln Ala Gly Pro Gly Thr Asp Ala Thr Ile Gln Ala Leu Leu Pro
 85 90 95

 His Leu Asp Lys Gly Asp Ile Leu Ile Asp Gly Gly Asn Thr Phe Tyr
 100 105 110

 Lys Asp Thr Ile Arg Arg Asn Glu Glu Leu Ala Asn Ser Gly Ile Asn

115	120	125
Phe Ile Gly Thr Gly Val Ser Gly Gly Glu Lys Gly Ala Leu Glu Gly 130 135 140		
Pro Ser Ile Met Pro Gly Gly Gln Lys Glu Ala Tyr Glu Leu Val Ala 145 150 155 160		
Asp Val Leu Glu Glu Ile Ser Ala Lys Ala Pro Glu Asp Gly Lys Pro 165 170 175		
Cys Val Thr Tyr Ile Gly Pro Asp Gly Ala Gly His Tyr Val Lys Met 180 185 190		
Val His Asn Gly Ile Glu Tyr Gly Asp Met Gln Leu Ile Ala Glu Ser 195 200 205		
Tyr Asp Leu Met Gln His Leu Leu Gly Leu Ser Ala Glu Asp Met Ala 210 215 220		
Glu Ile Phe Thr Glu Trp Asn Lys Gly Glu Leu Asp Ser Tyr Leu Ile 225 230 235 240		
Glu Ile Thr Ala Asp Ile Leu Ser Arg Lys Asp Asp Glu Gly Gln Asp 245 250 255		
Gly Pro Ile Val Asp Tyr Ile Leu Asp Ala Ala Gly Asn Lys Gly Thr 260 265 270		
Gly Lys Trp Thr Ser Gln Ser Ser Leu Asp Leu Gly Val Pro Leu Ser 275 280 285		
Leu Ile Thr Glu Ser Val Phe Ala Arg Tyr Ile Ser Thr Tyr Lys Glu 290 295 300		
Glu Arg Val His Ala Ser Lys Val Leu Pro Lys Pro Ala Ala Phe Asn 305 310 315 320		
Phe Glu Gly Asp Lys Ala Glu Leu Ile Glu Lys Ile Arg Gln Ala Leu 325 330 335		

Tyr Phe Ser Lys Ile Ile Ser Tyr Ala Gln Gly Phe Ala Gln Leu Arg
 340 345 350

Val Ala Ser Lys Glu Asn Asn Trp Asn Leu Pro Phe Ala Asp Ile Ala
 355 360 365

Ser Ile Trp Arg Asp Gly Cys Ile Ile Arg Ser Arg Phe Leu Gln Lys
 370 375 380

Ile Thr Asp Ala Tyr Asn Arg Asp Ala Asp Leu Ala Asn Leu Leu Leu
 385 390 395 400

Asp Glu Tyr Phe Leu Asp Val Thr Ala Lys Tyr Gln Gln Ala Val Arg
 405 410 415

Asp Ile Val Ala Leu Ala Val Gln Ala Gly Val Pro Val Pro Thr Phe
 420 425 430

Ser Ala Ala Ile Thr Tyr Phe Asp Ser Tyr Arg Ser Ala Asp Leu Pro
 435 440 445

Ala Asn Leu Ile Gln Ala Gln Arg Asp Tyr Phe Gly Ala His Thr Tyr
 450 455 460

Gln Arg Lys Asp Lys Glu Gly Thr Phe His Tyr Ser Trp Tyr Asp Glu
 465 470 475 480

Lys

<210> 630

<211> 144

<212> PRT

<213> Streptococcus pneumoniae

<400> 630

Met Asp Tyr Gln Arg Ile Asn Glu Tyr Leu Thr Ser Ile Phe Asn Asn
 1 5 10 15

Val Leu Val Ile Glu Glu Val Asn Leu Arg Gly Ser Arg Phe Lys Asp
 20 25 30

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Ile Ser Ile Lys Glu Met His Thr Ile Asp Val Ile Gly Lys Ala Pro
  35                      40                      45

Asp Val Thr Pro Ser Gln Val Ser Lys Glu Leu Met Val Thr Leu Gly
  50                      55                      60

Thr Val Thr Thr Ser Leu Asn Asn Leu Glu Arg Lys Gly Tyr Ile Glu
  65                      70                      75                      80

Arg Val Arg Ser Glu Gln Asp Arg Arg Val Val His Leu His Leu Thr
      85                      90                      95

Lys Lys Gly Arg Leu Ile His Arg Leu His Lys Arg Phe His Lys Ala
      100                      105                      110

Met Val Glu Lys Ile Ile Asp Gly Met Ser Glu Glu Glu Ile Ala Val
      115                      120                      125

Met Gly Lys Gly Leu Thr Asn Leu Tyr Gln Phe Leu Glu Asp Leu Lys
      130                      135                      140

<210> 631
<211> 74
<212> PRT
<213> Streptococcus pneumoniae

<400> 631

Met Ala Val Phe Glu Lys Val Gln Glu Ile Ile Val Glu Glu Leu Gly
  1                      5                      10                      15

Lys Asp Ala Ser Glu Val Thr Leu Glu Ser Thr Phe Asp Asp Leu Asp
      20                      25                      30

Ala Asp Ser Leu Asp Leu Phe Gln Val Ile Ser Glu Ile Glu Asp Ala
      35                      40                      45

Phe Asp Ile Gln Ile Glu Ala Glu Asn Asp Leu Lys Thr Val Gly Asp
      50                      55                      60

Leu Val Ala Tyr Val Glu Glu Gln Ala Lys
  65                      70

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<210> 632
 <211> 774
 <212> PRT
 <213> *Streptococcus pneumoniae*

 <400> 632

 Met Val Val Lys Thr Val Val Glu Ala Gln Asp Ile Phe Asp Lys Ala
 1 5 10 15

 Trp Glu Gly Phe Lys Gly Val Asp Trp Lys Glu Lys Ala Ser Val Ser
 20 25 30

 Arg Phe Val Gln Ala Asn Tyr Thr Pro Tyr Asp Gly Asp Glu Ser Phe
 35 40 45

 Leu Ala Gly Pro Thr Glu Arg Ser Leu His Ile Lys Lys Ile Val Glu
 50 55 60

 Glu Thr Lys Ala His Tyr Glu Glu Thr Arg Phe Pro Met Asp Thr Arg
 65 70 75 80

 Pro Thr Ser Ile Ala Asp Ile Pro Ala Gly Phe Ile Asp Lys Glu Asn
 85 90 95

 Glu Val Ile Phe Gly Ile Gln Asn Asp Glu Leu Phe Lys Leu Asn Phe
 100 105 110

 Met Pro Lys Gly Gly Ile Arg Met Ala Glu Thr Thr Leu Lys Glu Asn
 115 120 125

 Gly Tyr Glu Pro Asp Pro Ala Val His Glu Ile Phe Thr Lys Tyr Val
 130 135 140

 Thr Thr Val Asn Asp Gly Ile Phe Arg Ala Tyr Thr Ser Asn Ile Arg
 145 150 155 160

 Arg Ala Arg His Ala His Thr Val Thr Gly Leu Pro Asp Ala Tyr Ser
 165 170 175

 Arg Gly Arg Ile Ile Gly Val Tyr Ala Arg Leu Ala Leu Tyr Gly Ala
 180 185 190

Asp Tyr Leu Met Gln Glu Lys Val Asn Asp Trp Asn Ala Ile Lys Glu
 195 200 205

Ile Asp Glu Glu Thr Ile Arg Leu Arg Glu Glu Val Asn Leu Gln Tyr
 210 215 220

Gln Ala Leu Gln Gln Val Val Arg Leu Gly Asp Leu Tyr Gly Val Asp
 225 230 235 240

Val Arg Lys Pro Ala Met Asn Val Lys Glu Ala Ile Gln Trp Val Asn
 245 250 255

Ile Ala Phe Met Ala Val Cys Arg Val Ile Asn Gly Ala Ala Thr Ser
 260 265 270

Leu Gly Arg Val Pro Ile Val Leu Asp Ile Phe Ala Glu Arg Asp Leu
 275 280 285

Ala Arg Gly Thr Phe Thr Glu Ser Glu Ile Gln Glu Phe Val Asp Asp
 290 295 300

Phe Val Met Lys Leu Arg Thr Val Lys Phe Ala Arg Thr Lys Ala Tyr
 305 310 315 320

Asp Gln Leu Tyr Ser Gly Asp Pro Thr Phe Ile Thr Thr Ser Met Ala
 325 330 335

Gly Met Gly Asn Asp Gly Arg His Arg Val Thr Lys Met Asp Tyr Arg
 340 345 350

Phe Leu Asn Thr Leu Asp Asn Ile Gly Asn Ser Pro Glu Pro Asn Leu
 355 360 365

Thr Val Leu Trp Thr Asp Lys Leu Pro Tyr Asn Phe Arg Arg Tyr Cys
 370 375 380

Met His Met Ser His Lys His Ser Ser Ile Gln Tyr Glu Gly Val Thr
 385 390 395 400

Thr Met Ala Lys Asp Gly Tyr Gly Glu Met Ser Cys Ile Ser Cys Cys
 405 410 415

Val Ser Pro Leu Asp Pro Glu Asn Glu Glu Gln Arg His Asn Ile Gln
 420 425 430
 Tyr Phe Gly Ala Arg Val Asn Val Leu Lys Ala Leu Leu Thr Gly Leu
 435 440 445
 Asn Gly Gly Tyr Asp Asp Val His Lys Asp Tyr Lys Val Phe Asp Ile
 450 455 460
 Glu Pro Ile Arg Asp Glu Val Leu Glu Phe Glu Ser Val Lys Ala Asn
 465 470 475 480
 Phe Glu Lys Ser Leu Asp Trp Leu Thr Asp Thr Tyr Val Asp Ala Leu
 485 490 495
 Asn Ile Ile His Tyr Met Thr Asp Arg Tyr Asn Tyr Glu Ala Val Gln
 500 505 510
 Met Ala Phe Leu Pro Thr Lys Gln Arg Ala Asn Met Gly Phe Gly Ile
 515 520 525
 Cys Gly Phe Ala Asn Thr Val Asp Thr Leu Ser Ala Ile Lys Tyr Ala
 530 535 540
 Thr Val Lys Pro Ile Arg Asp Glu Asp Gly Tyr Ile Tyr Asp Tyr Glu
 545 550 555 560
 Thr Ile Gly Asp Tyr Pro Arg Trp Gly Glu Asp Asp Pro Arg Ser Asn
 565 570 575
 Glu Leu Ala Glu Trp Leu Ile Glu Ala Tyr Thr Thr Arg Leu Arg Ser
 580 585 590
 His Lys Leu Tyr Lys Asp Ala Glu Ala Thr Val Ser Leu Leu Thr Ile
 595 600 605
 Thr Ser Asn Val Ala Tyr Ser Lys Gln Thr Gly Asn Ser Pro Val His
 610 615 620
 Lys Gly Val Tyr Leu Asn Glu Asp Gly Ser Val Asn Leu Ser Lys Leu
 625 630 635 640

Glu Phe Phe Ser Pro Gly Ala Asn Pro Ser Asn Lys Ala Lys Gly Gly
645 650 655

Trp Leu Gln Asn Leu Asn Ser Leu Ser Ser Leu Asp Phe Ser Tyr Ala
660 665 670

Ala Asp Gly Ile Ser Leu Thr Thr Gln Val Ser Pro Arg Ala Leu Gly
675 680 685

Lys Thr Arg Asp Glu Gln Val Asp Asn Leu Val Thr Ile Leu Asp Gly
690 695 700

Tyr Phe Glu Asn Gly Gly Gln His Val Asn Leu Asn Val Met Asp Leu
705 710 715 720

Asn Asp Val Tyr Glu Lys Ile Met Ser Gly Glu Asp Val Ile Val Arg
725 730 735

Ile Ser Gly Tyr Cys Val Asn Thr Lys Tyr Leu Thr Pro Glu Gln Lys
740 745 750

Thr Glu Leu Thr Gln Arg Val Phe His Glu Val Leu Ser Met Asp Asp
755 760 765

Ala Leu Asp Ala Leu Ser
770

<210> 633

<211> 535

<212> PRT

<213> Streptococcus pneumoniae

<400> 633

Met Ser Thr Lys Tyr Ile Phe Val Thr Gly Gly Val Val Ser Ser Ile
1 5 10 15

Gly Lys Gly Ile Val Ala Ala Ser Leu Gly Arg Leu Leu Lys Asn Arg
20 25 30

Gly Leu Lys Val Thr Ile Gln Lys Phe Asp Pro Tyr Ile Asn Ile Asp
35 40 45

Pro Gly Thr Met Ser Pro Tyr Gln His Gly Glu Val Phe Val Thr Asp
 50 55 60

Asp Gly Ala Glu Thr Asp Leu Asp Leu Gly His Tyr Glu Arg Phe Ile
 65 70 75 80

Asp Ile Asn Leu Asn Lys Tyr Ser Asn Val Thr Thr Gly Lys Ile Tyr
 85 90 95

Ser Glu Val Leu Arg Lys Glu Arg Arg Gly Glu Tyr Leu Gly Ala Thr
 100 105 110

Val Gln Val Ile Pro His Ile Thr Asp Ala Leu Lys Glu Lys Ile Lys
 115 120 125

Arg Ala Ala Leu Thr Thr Asp Ser Asp Val Ile Ile Thr Glu Val Gly
 130 135 140

Gly Thr Val Gly Asp Ile Glu Ser Leu Pro Phe Leu Glu Ala Leu Arg
 145 150 155 160

Gln Met Lys Ala Asp Val Gly Ala Asp Asn Val Met Tyr Ile His Thr
 165 170 175

Thr Leu Leu Pro Tyr Leu Lys Ala Ala Gly Glu Met Lys Thr Lys Pro
 180 185 190

Thr Gln His Ser Val Lys Glu Leu Arg Gly Leu Gly Ile Gln Pro Asn
 195 200 205

Met Leu Val Ile Arg Thr Glu Glu Pro Ala Gly Gln Gly Ile Lys Asn
 210 215 220

Lys Leu Ala Gln Phe Cys Asp Val Ala Pro Glu Ala Val Ile Glu Ser
 225 230 235 240

Leu Asp Val Glu His Leu Tyr Gln Ile Pro Leu Asn Leu Gln Ala Gln
 245 250 255

Gly Met Asp Gln Ile Val Cys Asp His Leu Lys Leu Asp Ala Pro Ala

260	265	270
Ala Asp Met Thr Glu Trp Ser	Ala Met Val Asp Lys Val Met Asn Leu	
275	280	285
Lys Lys Gln Val Lys Ile Ser Leu Val Gly Lys Tyr Val Glu Leu Gln		
290	295	300
Asp Ala Tyr Ile Ser Val Val Glu Ala Leu Lys His Ser Gly Tyr Val		
305	310	315
Asn Asp Ala Glu Val Lys Ile Asn Trp Val Asn Ala Asn Asp Val Thr		
325	330	335
Ala Glu Asn Val Ala Glu Leu Leu Ser Asp Ala Asp Gly Ile Ile Val		
340	345	350
Pro Gly Gly Phe Gly Gln Arg Gly Thr Glu Gly Lys Ile Gln Ala Ile		
355	360	365
Arg Tyr Ala Arg Glu Asn Asp Val Pro Met Leu Gly Val Cys Leu Gly		
370	375	380
Met Gln Leu Thr Cys Ile Glu Phe Ala Arg His Val Leu Gly Leu Glu		
385	390	395
Gly Ala Asn Ser Ala Glu Leu Ala Pro Glu Thr Lys Tyr Pro Ile Ile		
405	410	415
Asp Ile Met Arg Asp Gln Ile Asp Ile Glu Asp Met Gly Gly Thr Leu		
420	425	430
Arg Leu Gly Leu Tyr Pro Ser Lys Leu Lys Arg Gly Ser Lys Ala Ala		
435	440	445
Ala Ala Tyr His Asn Gln Glu Val Val Gln Arg Arg His Arg His Arg		
450	455	460
Tyr Glu Phe Asn Asn Ala Phe Arg Glu Gln Phe Glu Ala Ala Gly Phe		
465	470	475
		480

Val Phe Ser Gly Val Ser Pro Asp Asn Arg Leu Val Glu Ile Val Glu
485 490 495

Ile Pro Glu Asn Lys Phe Phe Val Ala Cys Gln Tyr His Pro Glu Leu
500 505 510

Ser Ser Arg Pro Asn Arg Pro Glu Glu Leu Tyr Thr Ala Phe Val Thr
515 520 525

Ala Ala Val Glu Asn Ser Asn
530 535

<210> 634

<211> 378

<212> PRT

<213> Streptococcus pneumoniae

<400> 634

Met Asn Asn Thr Glu Phe Tyr Asp Arg Leu Gly Val Ser Lys Asn Ala
1 5 10 15

Ser Ala Asp Glu Ile Lys Lys Ala Tyr Arg Lys Leu Ser Lys Lys Tyr
20 25 30

His Pro Asp Ile Asn Lys Glu Pro Gly Ala Glu Asp Lys Tyr Lys Glu
35 40 45

Val Gln Glu Ala Tyr Glu Thr Leu Ser Asp Asp Gln Lys Arg Ala Ala
50 55 60

Tyr Asp Gln Tyr Gly Ala Ala Gly Ala Asn Gly Gly Phe Gly Gly Ala
65 70 75 80

Gly Gly Phe Gly Gly Phe Asn Gly Ala Gly Gly Phe Gly Gly Phe Glu
85 90 95

Asp Ile Phe Ser Ser Phe Phe Gly Gly Gly Ser Ser Arg Asn Pro
100 105 110

Asn Ala Pro Arg Gln Gly Asp Asp Leu Gln Tyr Arg Val Asn Leu Thr
115 120 125

Phe Glu Glu Ala Ile Phe Gly Thr Glu Lys Glu Val Lys Tyr His Arg
 130 135 140
 Glu Ala Gly Cys Arg Thr Cys Asn Gly Ser Gly Ala Lys Pro Gly Thr
 145 150 155 160
 Ser Pro Val Thr Cys Gly Arg Cys His Gly Ala Gly Val Ile Asn Val
 165 170 175
 Asp Thr Gln Thr Pro Leu Gly Met Met Arg Arg Gln Val Thr Cys Asp
 180 185 190
 Val Cys His Gly Arg Gly Lys Glu Ile Lys Tyr Pro Cys Thr Thr Cys
 195 200 205
 His Gly Thr Gly His Glu Lys Gln Ala His Ser Val His Val Lys Ile
 210 215 220
 Pro Ala Gly Val Glu Thr Gly Gln Gln Ile Arg Leu Ala Gly Gln Gly
 225 230 235 240
 Glu Ala Gly Phe Asn Gly Gly Pro Tyr Gly Asp Leu Tyr Val Val Val
 245 250 255
 Ser Val Glu Ala Ser Asp Lys Phe Glu Arg Glu Gly Thr Thr Ile Phe
 260 265 270
 Tyr Asn Leu Asn Leu Asn Phe Val Gln Ala Ala Leu Gly Asp Thr Val
 275 280 285
 Asp Ile Pro Thr Val His Gly Asp Val Glu Leu Val Ile Pro Glu Gly
 290 295 300
 Thr Gln Thr Gly Lys Lys Phe Arg Leu Arg Ser Lys Gly Ala Pro Ser
 305 310 315 320
 Leu Arg Gly Gly Ala Val Gly Asp Gln Tyr Val Thr Val Asn Val Val
 325 330 335
 Thr Pro Thr Gly Leu Asn Asp Arg Gln Lys Val Ala Leu Lys Glu Phe
 340 345 350

Ala Ala Ala Gly Asp Leu Lys Val Asn Pro Lys Lys Lys Gly Phe Phe
 355 360 365

Asp His Ile Lys Asp Ala Phe Asp Gly Glu
 370 375

<210> 635
 <211> 95
 <212> PRT
 <213> Streptococcus pneumoniae

<400> 635

Met Lys Leu Ser Asn Leu Leu Leu Phe Ala Gly Ala Ala Ala Gly Ser
 1 5 10 15

Tyr Leu Val Thr Lys Asn Arg Gln Thr Ile Thr Asp Glu Val Leu Asn
 20 25 30

Thr Thr Asp Arg Val Gln Ala Ile Lys Asp Asp Val Asp Ile Ile Gln
 35 40 45

Asn Ser Leu Gln Ile Ile Asn Gln Gln Lys Glu Leu Ile Lys Glu Tyr
 50 55 60

Gln Glu Asp Leu Thr Tyr Lys Phe Lys Val Leu Glu Lys Asp Ile Gln
 65 70 75 80

Thr Arg Leu Ala Val Ile Lys Glu Met Gln Gly Thr Glu Asp Lys
 85 90 95

<210> 636
 <211> 378
 <212> PRT
 <213> Streptococcus pneumoniae

<400> 636

Met Ser Lys Glu Met Leu Glu Ala Phe Arg Ile Leu Glu Glu Asp Lys
 1 5 10 15

Gly Ile Lys Lys Glu Asp Ile Ile Asp Ala Val Val Glu Ser Leu Arg
 20 25 30

Ser Ala Tyr Arg Arg Arg Tyr Gly Gln Ser Asp Ser Val Ala Ile Asp
 35 40 45

Phe Asn Glu Lys Thr Gly Asp Phe Thr Val Tyr Thr Val Arg Glu Val
 50 55 60

Val Asp Glu Val Phe Asp Ser Arg Leu Glu Ile Ser Leu Lys Asp Ala
 65 70 75 80

Leu Ala Ile Asn Ser Ala Tyr Glu Leu Gly Asp Lys Ile Lys Phe Glu
 85 90 95

Glu Ala Pro Ala Glu Phe Gly Arg Val Ala Ala Gln Ser Ala Lys Gln
 100 105 110

Thr Ile Met Glu Lys Met Arg Lys Gln Thr Arg Ala Ile Thr Tyr Asn
 115 120 125

Thr Tyr Lys Glu His Glu Gln Glu Ile Met Ser Gly Thr Val Glu Arg
 130 135 140

Phe Asp Asn Arg Phe Ile Tyr Val Asn Leu Gly Ser Ile Glu Ala Gln
 145 150 155 160

Leu Ser Lys Gln Asp Gln Ile Pro Gly Glu Val Phe Ala Ser His Asp
 165 170 175

Arg Ile Glu Val Tyr Val Tyr Lys Val Glu Asp Asn Pro Arg Gly Val
 180 185 190

Asn Val Phe Val Ser Arg Ser His Pro Glu Met Ile Lys Arg Leu Met
 195 200 205

Glu Gln Glu Ile Pro Glu Val Tyr Asp Gly Thr Val Glu Ile Met Ser
 210 215 220

Val Ala Arg Glu Ala Gly Asp Arg Thr Lys Val Ala Val Arg Ser His
 225 230 235 240

Asn Pro Asn Val Asp Ala Ile Gly Thr Ile Val Gly Arg Gly Gly Ala
 245 250 255

Asn Ile Lys Lys Ile Thr Ser Lys Phe His Pro Ala Arg Tyr Asp Ala
260 265 270

Lys Asn Asp Arg Met Val Pro Ile Glu Glu Asn Ile Asp Val Ile Glu
275 280 285

Trp Val Ala Asp Pro Ala Glu Phe Ile Tyr Asn Ala Ile Ala Pro Ala
290 295 300

Glu Val Asp Gln Val Ile Phe Asp Glu Asn Asp Ser Lys Arg Ala Leu
305 310 315 320

Val Val Val Pro Asp Asn Lys Leu Ser Leu Ala Ile Gly Arg Arg Gly
325 330 335

Gln Asn Val Arg Leu Ala Ala His Leu Thr Gly Tyr Arg Ile Asp Ile
340 345 350

Lys Ser Ala Ser Glu Phe Glu Ala Met Glu Asp Ala Ala Ser Val Glu
355 360 365

Leu Glu Val Glu Asn Asp Thr Val Glu Glu
370 375

<210> 637

<211> 958

<212> PRT

<213> Streptococcus pneumoniae

<400> 637

Met Ser Lys Lys Arg Leu Tyr Glu Ile Ala Lys Glu Leu Gly Lys Glu
1 5 10 15

Ser Lys Glu Val Val Ala Arg Ala Lys Glu Leu Gly Leu Asp Val Lys
20 25 30

Ser His Ser Ser Ser Val Glu Glu Ala Val Ala Ala Lys Ile Ala Ala
35 40 45

Ser Phe Lys Pro Ala Ala Ala Pro Lys Val Glu Ala Lys Pro Ala Ala
50 55 60

Pro Lys Val Ser Ala Glu Lys Lys Ala Glu Lys Ser Glu Pro Ala Lys
 65 70 75 80

Pro Ala Val Ala Lys Glu Glu Ala Lys Pro Ala Ala Pro Lys Ala Ser
 85 90 95

Ala Glu Lys Lys Ala Glu Lys Ser Glu Pro Val Lys Pro Ala Val Ala
 100 105 110

Lys Glu Glu Ala Lys Pro Ala Glu Pro Val Thr Pro Lys Thr Glu Lys
 115 120 125

Val Ala Ala Lys Pro Gln Ser Arg Asn Phe Lys Ala Glu Arg Glu Ala
 130 135 140

Arg Ala Lys Glu Gln Ala Glu Arg Arg Lys Gln Asn Lys Gly Asn Asn
 145 150 155 160

Arg Asp Gln Gln Gln Asn Gly Asn Arg Gln Lys Asn Asp Gly Arg Asn
 165 170 175

Gly Gly Lys Gln Gly Gln Ser Asn Arg Asp Asn Arg Arg Phe Asn Asp
 180 185 190

Gln Ala Lys Lys Gln Gln Gly Gln Gln Lys Arg Arg Asn Glu Arg Arg
 195 200 205

Gln Gln Glu Asp Lys Arg Ser Asn Gln Ala Ala Pro Arg Ile Asp Phe
 210 215 220

Lys Ala Arg Ala Ala Ala Leu Lys Ala Glu Gln Asn Ala Glu Tyr Ala
 225 230 235 240

Arg Ser Ser Glu Glu Arg Phe Lys Gln Tyr Gln Ala Ala Lys Glu Ala
 245 250 255

Leu Ala Gln Ala Asn Lys Arg Lys Glu Pro Glu Glu Ile Phe Glu Glu
 260 265 270

Ala Ala Lys Leu Ala Glu Gln Ala Gln Gln Val Gln Ala Val Val Glu
 275 280 285

Val Val Pro Glu Lys Lys Glu Pro Ala Val Asp Thr Arg Arg Lys Lys
 290 295 300
 Gln Ala Arg Pro Asp Lys Asn Arg Asp Asp Tyr Asp His Glu Glu Asp
 305 310 315 320
 Gly Pro Arg Lys Gln Gln Lys Asn Arg Ser Ser Gln Asn Gln Val Arg
 325 330 335
 Asn Gln Lys Asn Ser Asn Trp Asn Asn Asn Lys Lys Asn Lys Lys Gly
 340 345 350
 Asn Asn Lys Asn Asn Arg Asn Gln Thr Pro Lys Pro Val Thr Glu Arg
 355 360 365
 Lys Phe His Glu Leu Pro Thr Glu Phe Glu Tyr Thr Asp Gly Met Thr
 370 375 380
 Val Ala Glu Ile Ala Lys Arg Ile Lys Arg Glu Pro Ala Glu Ile Val
 385 390 395 400
 Lys Lys Leu Phe Met Met Gly Val Met Ala Thr Gln Asn Gln Ser Leu
 405 410 415
 Asp Gly Glu Thr Ile Glu Leu Leu Met Val Asp Tyr Gly Ile Glu Ala
 420 425 430
 Lys Gln Lys Val Glu Val Asp Asn Ala Asp Ile Glu Arg Phe Phe Val
 435 440 445
 Glu Asp Gly Tyr Leu Asn Glu Asp Glu Leu Val Glu Arg Pro Pro Val
 450 455 460
 Val Thr Ile Met Gly His Val Asp His Gly Lys Thr Thr Leu Leu Asp
 465 470 475 480
 Thr Leu Arg Asn Ser Arg Val Ala Thr Gly Glu Ala Gly Gly Ile Thr
 485 490 495
 Gln His Ile Gly Ala Tyr Gln Ile Val Glu Asn Gly Lys Lys Ile Thr

500	505	510
Phe Leu Asp Thr Pro Gly His Ala Ala Phe Thr Ser Met Arg Ala Arg 515	520	525
Gly Ala Ser Val Thr Asp Ile Thr Ile Leu Val Val Ala Ala Asp Asp 530	535	540
Gly Val Met Pro Gln Thr Ile Glu Ala Ile Asn His Ser Lys Ala Ala 545	550	555
Asn Val Pro Ile Ile Val Ala Ile Asn Lys Ile Asp Lys Pro Gly Ala 565	570	575
Asn Pro Glu Arg Val Ile Gly Glu Leu Ala Glu His Gly Val Met Ser 580	585	590
Thr Ala Trp Gly Gly Asp Ser Glu Phe Val Glu Ile Ser Ala Lys Phe 595	600	605
Asn Gln Asn Ile Glu Glu Leu Leu Glu Thr Val Leu Leu Val Ala Glu 610	615	620
Ile Gln Glu Leu Lys Ala Asp Pro Thr Val Arg Ala Ile Gly Thr Val 625	630	635
Ile Glu Ala Arg Leu Asp Lys Gly Lys Gly Ala Val Ala Thr Leu Leu 645	650	655
Val Gln Gln Gly Thr Leu Asn Val Gln Asp Pro Ile Val Val Gly Asn 660	665	670
Thr Phe Gly Arg Val Arg Ala Met Thr Asn Asp Leu Gly Arg Arg Val 675	680	685
Lys Val Ala Gly Pro Ser Thr Pro Val Ser Ile Thr Gly Leu Asn Glu 690	695	700
Ala Pro Met Ala Gly Asp His Phe Ala Val Tyr Glu Asp Glu Lys Ser 705	710	715
		720

Ala Arg Ala Ala Gly Glu Glu Arg Ala Lys Arg Ala Leu Met Lys Gln
725 730 735

Arg Gln Ala Thr Gln Arg Val Ser Leu Glu Asn Leu Phe Asp Thr Leu
740 745 750

Lys Ala Gly Glu Leu Lys Ser Val Asn Val Ile Ile Lys Ala Asp Val
755 760 765

Gln Gly Ser Val Glu Ala Leu Ser Ala Ser Leu Gln Lys Ile Asp Val
770 775 780

Glu Gly Val Lys Val Thr Ile Val His Ser Ala Val Gly Ala Ile Asn
785 790 795 800

Glu Ser Asp Val Thr Leu Ala Glu Ala Ser Asn Ala Phe Ile Val Gly
805 810 815

Phe Asn Val Arg Pro Thr Pro Gln Ala Arg Gln Gln Ala Glu Ala Asp
820 825 830

Asp Val Glu Ile Arg Leu His Ser Ile Ile Tyr Lys Val Ile Glu Glu
835 840 845

Met Glu Glu Ala Met Lys Gly Met Leu Asp Pro Glu Phe Glu Glu Lys
850 855 860

Val Ile Gly Glu Ala Val Ile Arg Glu Thr Phe Lys Val Ser Lys Val
865 870 875 880

Gly Thr Ile Gly Gly Phe Met Val Ile Asn Gly Lys Val Ala Arg Asp
885 890 895

Ser Lys Val Arg Val Ile Arg Asp Gly Val Val Ile Tyr Asp Gly Glu
900 905 910

Leu Ala Ser Leu Lys His Tyr Lys Asp Asp Val Lys Glu Val Thr Asn
915 920 925

Gly Arg Glu Gly Gly Leu Met Ile Asp Gly Tyr Asn Asp Ile Lys Met
930 935 940

Asp Asp Val Ile Glu Ala Tyr Val Met Glu Glu Ile Lys Arg
 945 950 955

<210> 638
 <211> 293
 <212> PRT
 <213> Streptococcus pneumoniae

<400> 638

Met Ala Ile Val Ser Ala Glu Lys Phe Val Gln Ala Ala Arg Asp Asn
 1 5 10 15

Gly Tyr Ala Val Gly Gly Phe Asn Thr Asn Asn Leu Glu Trp Thr Gln
 20 25 30

Ala Ile Leu Arg Ala Ala Glu Ala Lys Lys Ala Pro Val Leu Ile Gln
 35 40 45

Thr Ser Met Gly Ala Ala Lys Tyr Met Gly Gly Tyr Lys Val Ala Arg
 50 55 60

Asn Leu Ile Ala Asn Leu Val Glu Ser Met Gly Ile Thr Val Pro Val
 65 70 75 80

Ala Ile His Leu Asp His Gly His Tyr Glu Asp Ala Leu Glu Cys Ile
 85 90 95

Glu Val Gly Tyr Thr Ser Ile Met Phe Asp Gly Ser His Leu Pro Val
 100 105 110

Glu Glu Asn Leu Lys Leu Ala Lys Glu Val Val Glu Lys Ala His Ala
 115 120 125

Lys Gly Ile Ser Val Glu Ala Glu Val Gly Thr Ile Gly Gly Glu Glu
 130 135 140

Asp Gly Ile Ile Gly Lys Gly Glu Leu Ala Pro Ile Glu Asp Ala Lys
 145 150 155 160

Ala Met Val Glu Thr Gly Ile Asp Phe Leu Ala Ala Gly Ile Gly Asn
 165 170 175

Ile His Gly Pro Tyr Pro Val Asn Trp Glu Gly Leu Asp Leu Asp His
 180 185 190

Leu Gln Lys Leu Thr Glu Ala Leu Pro Gly Phe Pro Ile Val Leu His
 195 200 205

Gly Gly Ser Gly Ile Pro Asp Glu Gln Ile Gln Ala Ala Ile Lys Leu
 210 215 220

Gly Val Ala Lys Val Asn Val Asn Thr Glu Cys Gln Ile Ala Phe Ala
 225 230 235 240

Asn Ala Thr Arg Lys Phe Ala Arg Asp Tyr Glu Ala Asn Glu Ala Glu
 245 250 255

Tyr Asp Lys Lys Lys Leu Phe Asp Pro Arg Lys Phe Leu Ala Asp Gly
 260 265 270

Val Lys Ala Ile Gln Ala Ser Val Glu Glu Arg Ile Asp Val Phe Gly
 275 280 285

Ser Glu Gly Lys Ala
 290

<210> 639

<211> 141

<212> PRT

<213> Streptococcus pneumoniae

<400> 639

Met Ala Lys Lys Val Glu Lys Leu Val Lys Leu Gln Ile Pro Ala Gly
 1 5 10 15

Lys Ala Thr Pro Ala Pro Pro Val Gly Pro Ala Leu Gly Gln Ala Gly
 20 25 30

Ile Asn Ile Met Gly Phe Thr Lys Glu Phe Asn Ala Arg Thr Ala Asp
 35 40 45

Gln Ala Gly Met Ile Ile Pro Val Val Ile Ser Val Tyr Glu Asp Lys
 50 55 60

Ser Phe Thr Phe Val Thr Lys Thr Pro Pro Ala Ala Val Leu Leu Lys
65 70 75 80

Lys Ala Ala Gly Val Glu Lys Gly Ser Gly Thr Pro Asn Lys Thr Lys
85 90 95

Val Ala Thr Val Thr Arg Ala Gln Val Gln Glu Ile Ala Glu Thr Lys
100 105 110

Met Pro Asp Leu Asn Ala Ala Asn Val Glu Ser Ala Met Arg Met Ile
115 120 125

Glu Gly Thr Ala Arg Ser Met Gly Phe Thr Val Val Asp
130 135 140

<210> 640
<211> 229
<212> PRT
<213> Streptococcus pneumoniae

<400> 640

Met Ala Lys Lys Ser Lys Gln Leu Arg Ala Ala Leu Glu Lys Ile Asp
1 5 10 15

Ser Thr Lys Ala Tyr Ser Val Glu Glu Ala Val Ala Leu Ala Lys Glu
20 25 30

Thr Asn Phe Ala Lys Phe Asp Ala Thr Val Glu Val Ala Tyr Asn Leu
35 40 45

Asn Ile Asp Val Lys Lys Ala Asp Gln Gln Ile Arg Gly Ala Met Val
50 55 60

Leu Pro Asn Gly Thr Gly Lys Thr Ser Arg Val Leu Val Phe Ala Arg
65 70 75 80

Gly Ala Lys Ala Glu Glu Ala Lys Ala Ala Gly Ala Asp Phe Val Gly
85 90 95

Glu Asp Asp Leu Val Ala Lys Ile Asn Asp Gly Trp Leu Asp Phe Asp
100 105 110

Val Val Ile Ala Thr Pro Asp Met Met Ala Leu Val Gly Arg Leu Gly
 115 120 125

Arg Val Leu Gly Pro Arg Asn Leu Met Pro Asn Pro Lys Thr Gly Thr
 130 135 140

Val Thr Met Asp Val Ala Lys Ala Val Glu Glu Ser Lys Gly Gly Lys
 145 150 155 160

Ile Thr Tyr Arg Ala Asp Arg Ala Gly Asn Val Gln Ala Ile Ile Gly
 165 170 175

Lys Val Ser Phe Glu Ala Glu Lys Leu Val Glu Asn Phe Lys Ala Phe
 180 185 190

Asn Glu Thr Ile Gln Lys Ala Lys Pro Ala Thr Ala Lys Gly Thr Tyr
 195 200 205

Val Thr Asn Leu Thr Ile Thr Thr Thr Gln Gly Val Gly Ile Lys Val
 210 215 220

Asp Val Asn Ser Leu
 225

<210> 641
 <211> 620
 <212> PRT
 <213> Streptococcus pneumoniae

<400> 641

Met Asn Ile Ile Glu Glu Ile Met Thr Lys Leu Arg Glu Asp Ile Arg
 1 5 10 15

Asn Ile Ala Ile Ile Ala His Val Asp His Gly Lys Thr Thr Leu Val
 20 25 30

Asp Glu Leu Leu Lys Gln Ser Glu Thr Leu Asp Ala Arg Thr Glu Leu
 35 40 45

Ala Glu Arg Ala Met Asp Ser Asn Asp Ile Glu Lys Glu Arg Gly Ile
 50 55 60

Thr Ile Leu Ala Lys Asn Thr Ala Val Ala Tyr Asn Gly Thr Arg Ile
 65 70 75 80

Asn Ile Met Asp Thr Pro Gly His Ala Asp Phe Gly Gly Glu Val Glu
 85 90 95

Arg Ile Met Lys Met Val Asp Gly Val Val Leu Val Val Asp Ala Tyr
 100 105 110

Glu Gly Thr Met Pro Gln Thr Arg Phe Val Leu Lys Lys Ala Leu Glu
 115 120 125

Gln Asp Leu Val Pro Ile Val Val Val Asn Lys Ile Asp Lys Pro Ser
 130 135 140

Ala Arg Pro Ala Glu Val Val Asp Glu Val Leu Glu Leu Phe Ile Glu
 145 150 155 160

Leu Gly Ala Asp Asp Asp Gln Leu Asp Phe Pro Val Val Tyr Ala Ser
 165 170 175

Ala Ile Asn Gly Thr Ser Ser Leu Ser Asp Asp Pro Ala Asp Gln Glu
 180 185 190

Ala Thr Met Ala Pro Ile Phe Asp Thr Ile Ile Asp His Ile Pro Ala
 195 200 205

Pro Val Asp Asn Ser Asp Glu Pro Leu Gln Phe Gln Val Ser Leu Leu
 210 215 220

Asp Tyr Asn Asp Phe Val Gly Arg Ile Gly Ile Gly Arg Val Phe Arg
 225 230 235 240

Gly Thr Val Lys Val Gly Asp Gln Val Thr Leu Ser Lys Leu Asp Gly
 245 250 255

Thr Thr Lys Asn Phe Arg Val Thr Lys Leu Phe Gly Phe Phe Gly Leu
 260 265 270

Glu Arg Arg Glu Ile Gln Glu Ala Lys Ala Gly Asp Leu Ile Ala Val
 275 280 285